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9 S24 AND (S1 OR HEMOGLOBIN?)

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DIALOG(R)File 5:Biosis Previews(R)
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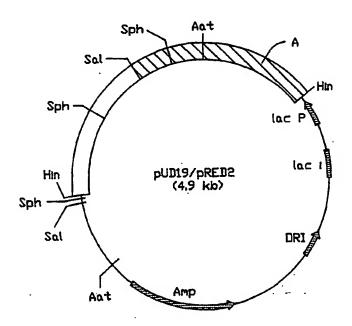
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(54) Title: ENHANCEMENT OF CELL GROWTH BY EXPRESSION OF A CLONED HEMOGLOBIN GENE



(57) Abstract

The invention related to a gene promoter/regulator (Region A in Fig. 1) which is useful in subjecting expression of protein or polypeptides in a host cell to selective regulation by external control. In particular, this regulation may be accomplished to different extends by the level of dissolved oxygen, presence of cAMP-CAP, and/or a complex nitrogen source in the culture or environment. The invention also relates to a method of improving production of proteins in a host by coproduction of hemoglobin in the host.

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ENHANCEMENT OF CELL GROWTH BY EXPRESSION OF A CLONED HEMOGLOBIN GENE

This is a continuation-in-part of Serial No. 342,451, filed January 24, 1989 as PCT application No. PCT

5 US88-03745, which is a continuation-in-part of Serial No. 113,014 filed October 23, 1987 and Serial No. 151,526, filed February 2, 1988.

TECHNICAL FIELD

This invention relates to the production of proteins
and polypeptides, and to production of oxygen-binding
proteins, particularly hemoglobins, and to
enhancement of the growth and product synthesis
characteristics of aerobic organisms in environments
with sufficient as well as reduced or low levels of
oxygen.

This invention relates generally to the use of recombinant DNA technology to direct or otherwise control gene expression in cultured cells, and more particularly, to methods and materials useful in subjecting the transcription and translation of DNA sequences to selective regulation by external control.

BACKGROUND ART

Globins such as hemoglobin and myoglobin are hemecontaining oxygen carriers. By reversibly binding to oxygen in the presence of high oxygen 5 concentrations and releasing it in regions or at times of low concentrations, these proteins considerably enhance the oxygen uptake rate of multicellular organisms over that allowed by mere passive diffusion. In unicellular organisms it is 10 generally believed that the oxygen uptake rate is principally limited by the rate of transfer of dissolved oxygen in the environment or growth medium to the exterior cell surface. However, closer examination of cell structure reveals several 15 potential diffusional barriers between environmental oxygen and the cytochromes where the oxygen finally undergoes reaction. For example, in gram negative bacteria, where the cytochromes are attached to the inside of the plasma membrane, the diffusing oxygen 20 needs to cross transport barriers such as the cell wall, the outer membrane, the periplasmic space and the inner membrane before accepting electrons from metabolic reactions. In unicellular eucaryotes, where oxidative phosphorylation takes place in the 25 mitochondria, there are further diffusional resistances. Small neutral molecules like oxygen are assumed to passively diffuse across these barriers; however, these barriers make a non-trivial contribution to the overall resistance to mass 30 transfer to the actual reaction site and thus could

Physiological effects on growth due to depletion in dissolved oxygen levels has been demonstrated in the case of several organisms, including <u>Escherichia</u>

35 <u>coli</u>, <u>Saccharomyces cerevisiae</u>, <u>Pseudomonas</u> strains, and <u>Alcaligenes eutrophus</u>. In <u>E. coli</u> for example,

be of significance under oxygen-limited conditions.

which has a very high affinity cytochrome, changes in dissolved oxygen tension leads to differential regulation of terminal oxidases, resulting in a decrease in the number of protons expelled per NADH mecule oxidized during aerobic respiration and, consequently, a possible adverse change in the stoichiometry of ATP biosynthesis. (Kranz et al., Journal of Bacteriology 158:1191-1194, 1984; Ingraham et al., Growth of the bacterial cell, Sinauer Associates, Inc. 1983, p. 147, both specifically incorporated herein.)

In addition to the respiratory oxygen requirement of aerobic organisms, oxygen-binding proteins have other potential applications as well, including, for example, the enhancement of particular oxidative transformations such as steroid conversions, vinegar production, biological waste treatment or enzymatic degradations, and in some steps in brewing or making distilled and fermented foods and beverages.

- The filamentous bacterium, <u>Vitreoscilla</u>, a member of the Beggiatoa family, is a strict aerobe that is found in oxygen-poor environments such as stagnant ponds and decaying vegetable matter. Growth of the bacterium under hypoxic conditions results in a
- 25 several-fold induction of synthesis of a homodimeric soluble heme protein (subunit MW 15,775) (Boerman et al., Control of heme content in Vitreoscilla by oxygen, Journal of General Applied Microbiology 28:35-42, 1982) which has a remarkable spectral
- 30 (Webster, et al., Reduced nicotinamide adenine dinucleotide cytochrome o reductase associated with cytochrome o purified from Vitreoscilla, Journal of Biological Chemistry 249:4257-4260, 1974), structural (Wakabayashi, et al., Primary sequences of a dimeric
- 35 bacterial hemoglobin from Vitreoscilla, Nature

322:481-483, 1986), and kinetic (Orii, et al.,

Photodissociation of oxygenated cytochrome o(xs)

(Vitreoscilla) and kinetic studies of reassociation,

Journal of Biol gical Chemistry 261:2978-2986, 1986)

5 homology with eucaryotic hemoglobins, and which is

probably a true bacterial hemoglobin.

This protein was previously thought to be a cytochrome o, and it has been suggested to function in oxygen storage. However, biochemical 10 discrepancies (Webster, et al., Oxygenated cytochrome Q, Journal of Biological Chemistry 252:1834-1836, 1977) as well as the subsequent discovery of the true membrane-bound cytochromes o and d (DeMaio, et al., Spectral evidence for the existence of a second 15 cytochrome o in whole cells of Vitreoscilla, Journal of Biological Chemistry 258:13768-13771, 1983; Webster et al., Federation Proceeding 44:678, 1985) led to further investigations of its spectral properties (Choc et al., Oxygenated intermediate and 20 carbonyl species of cytochrome o (Vitreoscilla), Journal of Biological Chemistry 257: 865-869, 1982; Orii et al., supra.) and the eventual determination of its probable amino acid sequences and partial homology with known hemoglobin sequences (23).

Although these articles disclose the conservation of most features characteristic of eucaryotic hemoglobins, and discuss, to some extent, the role or potential role it probably plays in oxygen utilization, none of these researchers had previously been able to isolate a portable DNA sequence capable of directing intracellular production of this bacterial hemoglobin or to create a recombinant-DNA method for its production. Additionally, there has been no published proof of any oxygen transport or other kinetic function for this protein in

<u>Vitreoscilla</u>, or any suggestion in the literature of any benefit from the introduction of a bacterial hemoglobin in heterologous organisms. Moreover, there has been no suggestion that such an oxygen-binding protein would have a far-reaching range of applications.

Surprisingly, the present inventors have discovered a portable DNA sequence capable of directing the recombinant-DNA synthesis of a bacterial hemoglobin. 10 The hemoglobin of the present invention, prepared by the recombinant-DNA methods set forth herein, will enable increased research into the growth of organisms in oxygen-poor environments. In addition, the oxygen-binding proteins of the present invention 15 are useful in enhancing oxygen supply to cells or in other oxygen-utilizing processes, and for binding and separating oxygen from other fluids or gases. Furthermore, the oxygen-binding proteins of this invention are capable of increasing production of 20 cells, or of proteins or metabolites normally made by a cell, or of natural or unnatural metabolites and proteins expressed in a cell via genetic manipulation. The proteins of this invention are also useful as selective markers in recombinant-DNA 25 work, and have applications as diverse as enhancing certain oxygen-requiring steps in fermentation, enzymatic degradation, toxic chemical waste treatment, brewing, and particular oxidative reactions and transformations.

This invention also relates to certain DNA sequences which usually precede a gene in a DNA polymer and which provide a site for initiation of the transcription of that gene into mRNA. These are referred to as "promoter" sequences. Other DNA or RNA sequences, also usually but not necessarily

"upstream" of a structural gene, bind proteins that determine the frequency or rate of transcription and/or translation initiation. These ther sequences, including attenuators, enhancers, operators and the like, are referred to as "regulator" sequences. Thus, sequences which operate to determine whether the transcription and eventual expression of a gene will take place are collectively referred to as "promoter/regulator" DNA sequences.

10 The promoter/regulator sequences of genes are susceptible to enormous structural and functional variation, and in general, serve to regulate gene transcription in response to chemical and, sometimes, physical environmental conditions in and around the 15 cell. Several generalized models for the action of promoter/regulator operation in gene transcription have been proposed. One model utilizes a repressor gene and a regulator sequence or operator sequence near the promoter of another gene. According to this 20 model, transcription of the repressor sequence results in expression of a repressor protein which selectively binds to the operator sequence to effectively preclude transcription of the selected gene. An environmental signal, such as the increased 25 concentration of a chemical acted upon by the protein product of the gene in question, may operatively inactivate the repressor protein, blocking its ability to bind to the operator sequence in a way which would interrupt transcription of the gene. 30 Increased concentrations of a substrate could be seen

Another generalized model of operation of promoter/regulator sequences in the regulation of gene transcription suggests formation of an initially

as operating to induce synthesis of the protein which

catalyzes its breakdown.

inactive form of repressor protein by the repressor DNA sequence. Such inactive form could not bind to an operator DNA sequence and disrupt selected gene transcription until it is combined with some other

substance present in the cell, such as a compound which is the product of a reaction catalyzed by the protein coded for by the selected gene. Increased concentrations of such a reaction product in the cell would thus operate to repress the potential

overproduction of proteins responsible for the product's synthesis. In these examples, the regulator protein functions to inhibit transcription. Other regulatory proteins have been described which potentiate or activate transcription of specific DNA

15 sequences. Thus, there can be both positive and negative control proteins and corresponding regulatory DNA sequences.

Regulation of gene expression can also occur at the level of translation. For example, a regulator

molecule could bind to a particular site on the messenger RNA, thus inhibiting or blocking translation.

Much of the genetic engineering activity to date has been oriented toward stably incorporating foreign DNA into cells, to provide not only a source of multiple copies of selected genes, but the large scale transcription and expression of commercially significant gene products.

The lactose ("lac") promoter/operator systems have

been commonly used, for they are very controllable
through the mode of action of the operator. When the
operator is repressed, the DNA dependent RNA
polymerase is completely prevented from binding and
initiating transcription, thus effectively blocking

promoter operability. This system can be derepressed by induction following the addition of a known inducer, such as isopropyl-beta-D-thiogalactoside (IPTG). The inducer causes the repressor protein to fall away so the RNA polymerase can function.

Cells transformed with plasmids carrying the lac promoter/operator system can be permitted to grow up to maximal density while in the repressed state through the omission of an inducer, such as IPTG, from the media. When a high level of cell density is achieved, the system can be derepressed by addition of inducer. The promoter is then free to initiate transcription and thus obtain expression of the gene products at yields commensurate with the promoter strength. However, certain of these inducible promoter systems are relatively weak and commercial or research productions using such systems do not urge the cell to generate maximum output.

In response to the need for microbial expression 20 vehicles capable of producing desired products in higher yield, the tryptophan ("trp") promoter/ operator system has become widely used. This system is one of several known systems with at least three times the strength of the lac promoter. However, it 25 has the disadvantage of less promoter control. trp promoter is not inducible in the way the lac promoter is, namely, the bound repressor is not removed by induction. Instead, the system operates on a sort of feedback loop as described above. A 30 system was devised whereby the attenuator region of the trp promoter/operator system was removed, with the resultant transformed cells being grown in tryptophan-rich media. This provided sufficient tryptophan to essentially completely repress the 35 operator so that cell growth could proceed

uninhibited by premature expression of any desired foreign prot ins. When the culture reached appropriat growth levels, no additional tryptophan was supplied, resulting in mild tryptophan

5 limitation, and, accordingly, derepression of the promoter with resultant expression of the desired protein gene insert. In application, this system has several disadvantages. For example, it is necessary to maintain high levels of tryptophan in the growth media to completely repress the promoter, and to permit the medium to become completely exhausted of tryptophan following full growth of the culture.

A hybrid system has been developed from the tryptophan and lactose promoter, wherein both promoters can be repressed by the lac repressor and both can be derepressed with IPTG. See De Boer et al., The tac promoter: A functional hybrid derived from the trp and lac promoters, Proc. Natl. Acad. Sci. USA, 80: 21-25, 1983. This system shares a disadvantage with the two discussed above, namely the required introduction of additional agents to a normal growth medium.

Another regulator/promoter system commonly used for expression of cloned proteins in <u>E. coli</u> is based on the P_L promoter system from phage lambda. See Bernard and Helsinki, Methods in Enzymology, 68:482-492, 1979; <u>Use of Lambda Phage Promoter PL to Promote Gene Expression In Hybrid Plasmid Cloning Vehicles</u>. Induction of this promoter requires increase of culture temperature from 30°C to 42°C. This system has the disadvantages of suboptimal growth rates at 30°C prior to induction and upsetting of cell metabolism by the temperature shift. Temperature shift effects on metabolism are discussed, for example, by Neidhart, <u>et. al.</u>, <u>The Genetics and</u>

Regulation Of Heat-Shock Proteins, Annual Reviews of Genetics, 18:295-329, 1984.

There has been a need in the art for an economical, simple, highly controllable and efficient

5 promoter/regulator system for subjecting the transcription of DNA sequences to selective regulation by external control at constant temperature. The present inventors have discovered such an expression system, which can switch from low to very high expression activity upon reduction of dissolved oxygen concentration in the medium. This reduction in dissolved oxygen (DO) level is easily implemented at high cell densities without the need for addition of any chemical to the growth medium to induce gene expression.

In addition, other modes of regulation independent of or in conjunction with control of DO have been discovered.

DISCLOSURE OF THE INVENTION

The present invention relates to portable DNA sequences capable of directing intracellular production of proteins and polypeptides, including oxygen-binding proteins, particularly hemoglobins. The present invention also relates to vectors containing these portable DNA sequences.

One object of the present invention is to provide an active (<u>i.e.</u>, oxygen-transporting form) bacterial hemoglobin protein, which can be produced in sufficient quantities and purities to provide economical pharmaceutical, laboratory or industrial compositions which possess oxygen-binding activity.

An additional object of the present invention is to provid a recombinant-DNA method for the production of these oxygen-binding proteins. To facilitate the recombinant-DNA synthesis of these oxygen-binding 5 proteins, it is a further object of the present invention to provide portable DNA sequences capable of directing intracellular production of oxygenbinding proteins. It is also an object of the present invention to provide cloning vectors 10 containing these portable sequences. These vectors are capable of being used in recombinant systems to enhance the growth characteristics of organisms, and to produce useful quantities of oxygen-binding proteins. Augmented by intracellular synthesis of 15 oxygen-binding proteins, product formation may also be enhanced.

The present invention also provides novel methods and materials for subjecting DNA sequences of living microorganisms to external regulation which is

20 dependent upon availability of oxygen in the environment. Particularly, it relates to promoter/regulators, a recombinant-DNA method of producing same, and to portable DNA sequences capable of directing the translation and transcription

25 initiation and control of the expression of desired gene products.

Thus, another object of the present invention is to provide for the control of expression of any selected chromosomal or extrachromosomal gene or DNA sequence through the incorporation of a promoter/regulator DNA sequence which is functionally responsive to environmental variations in the concentration of oxygen. The invention is thus broadly applicable to a variety of aerobic or slightly aerobic procedures for controlling genetic processes, ranging from the

alteration of existing regulation of endogenous genes in prokaryotic and eucaryotic cells to securing selectiv, differential regulation of expression of selected exogenous or foreign genes stably incorporated in host cells.

Still another object of the present invention is to provide a promoter/regulator as described above, which can be produced in sufficient quantities and purities to provide their economical pharmaceutical, laboratory or industrial use.

Additional objects and advantages of the invention will be set forth in part in the description which follows, and in part will be obvious from the description or may be learned from practice of the invention. The objects and advantages may be realized and attained by means of the instrumentalities and combination particularly pointed out in the appended claims.

To achieve the objects and in accordance with the

20 purposes of the present invention, promoter/
regulators are also set forth. To further achieve
the objects and in accordance with the purposes of
the present invention, as embodied and broadly
described herein, portable DNA sequences for these

25 promoter/regulators are provided. Particularly
preferred promoter/regulator DNA sequences for
use in the practice of the present invention are
derived from the filamentous bacterium <u>Vitreoscilla</u>.
Portable nucleotide sequences are provided for these

30 promoter/regulators. The portable sequences may be
either synthetic sequences or restriction fragments
("natural" DNA sequences).

To facilitate identification and isolation of natural DNA sequences for use in the present invention, the inventors have developed a <u>Vitreoscilla</u> genomic library. This library contains the genetic information capabl of directing a cell to synthesize the hemoglobin of the present invention. Other

the hemoglobin of the present invention. Other natural DNA sequences which may be used in the recombinant DNA methods set forth herein may be isolated from other genomic libraries.

10 Additionally, portable DNA sequences useful in the processes of the present invention may be synthetically created. These synthetic DNA sequences may be prepared by polynucleotide synthesis and sequencing techniques known to those of ordinary skill in the art.

Additionally, to achieve the objects and in accordance with the purposes of the present invention, a recombinant-DNA method is disclosed which results in manufacture by a host cell or 20 microorganism of oxygen-binding proteins or other cloned proteins or polypeptides using the portable DNA sequences referred to above.

Additionally, to achieve the objects and in accordance with the purposes of the present invention, recombinant-DNA methods are disclosed which subject to external control the translation and transcription of gene products by a host cell or microorganism using the portable DNA sequences referred to above.

30 Processes of the invention include methods for subjecting the expression of a selected DNA sequence in a living cell or virus to regulation by oxygen level through the site-specific insertion of

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promoter/regulator DNA sequences responsive thereto. Also disclosed are improvements in prior methods for securing expression of a selected "foreign" or exogenous sequence in a host microorganism wherein 5 the DNA sequence is stably incorporated as chromosomal or extrachromosomal constituent of the host. Such improvements comprise fusing to the selected DNA sequence a promoter/regulator DNA sequence capable of selectively promoting or 10 inhibiting expression of the selected DNA in response to variations in environmental concentration of oxygen. It has now been found that expression of oxygen-binding proteins, particularly in oxygen-poor environments, can improve intracellular production of 15 other proteins. Thus, a host cell containing a first expression vector (extrachromosomal) for expression of a designated protein may be modified to enhance production of that protein by introducing into the host a second expression vector 20 (which may or may not be part of the first expression vector) for expression of an oxygen-binding protein. In an oxygen-poor environment, the second vector improves the cell productivity, and thus the production of the designated protein.

25 It has further been found that there are modes of regulating the activity of the Vitreoscilla hemoglobin promoter/regulator other than by environmental oxygen concentration (DO). During the exponential growth phase of a host cell culture, it 30 has been found that addition of a complex nitrogen source, such as yeast extract, to the culture allows suppression of the Vitreoscilla hemoglobin promoter/regulator independently of environmental oxygen concentration. It has also been found that 35 the presence of the cAMP-CAP complex is important to the activity of the Vitreoscilla hemoglobin

promoter/regulator, thus manipulation of the activity crp gene in a host cell is another method of regulation. For example, using a crp mutant host, by direct addition of CAP t the culture the activity of the <u>Vitreoscilla</u> promoter/regulator may be regulated.

It is understood that both the foregoing general description and the following detailed description are exemplary and explanatory only and are not restrictive of the invention, as claimed.

- 10 The accompanying drawing, which is incorporated in and constitutes a part of this specification, illustrates one embodiment of the invention and, together with the description, serves to explain the principles of the invention.
- 15 <u>BRIEF DESCRIPTION OF THE DRAWINGS</u>
 Figure 1 is a partial restriction map of the plasmid pUC19/pRED2.

Reference will now be made in detail to the presently preferred embodiments of the invention, which, together with the drawing and the following examples, serve to explain the principles of the invention.

It must be understood that the present inventors have prepared an oxygen-binding protein and its natural promoter/regulator by recombinant DNA methods. While some methods for the production and use of these recombinant products are described below, the end use of these products alone is within the scope of the present invention.

30 As noted above, the present invention relates in part to portable DNA sequences capable of directing

intracellular production of oxygen-binding proteins in a variety of host cells and host microorganisms. "Portable DNA sequence" in this context is intended to refer either to a synthetically produced 5 nucleotide sequence or to a restriction fragment of a naturally occurring DNA sequence. For purposes of this specification, "oxygen-binding protein" is intended to mean a protein with a primary structure as defined by the codons present in the 10 deoxyribonucleic acid sequence which directs intracellular production of the amino acid sequence, and which may or may not include post-translational modifications. It is contemplated that such posttranslational modifications include, for example, 15 association with a heme prosthetic group. further intended that the term "oxygen-binding protein" refers to either the form of the protein as would be excreted from a cell or as it may be present in the cell from which it was not excreted.

In a preferred embodiment, the portable DNA sequences are capable of directing intracellular production of hemoglobin. In a particularly preferred embodiment, the portable DNA sequences are capable of directing intracellular production of a hemoglobin biologically equivalent to that previously isolated from the filamentous bacterium, Vitreoscilla. By "biologically equivalent", as used herein, it is meant that a protein, produced using a portable DNA sequence of the present invention, is capable of binding oxygen in the same fashion, but not necessarily to the same degree, as the homodimeric soluble heme protein (subunit MW 15,775) isolable from Vitreoscilla.

As noted above, the present invention also relates in part to portable DNA sequences which contain

promoter/regulators which are capable of directing intracellular expression f endogenous or exogenous gene products, in a variety of host cells and h st microorganisms. "Portable DNA sequence" and

5 "promoter/regulator" in this context are intended to refer either to a synthetically produced nucleotide sequence or to a restriction fragment of a naturally occurring DNA sequence.

The portable DNA sequences of the present invention

10 may also include DNA sequences downstream from a
promoter/regulator which code for at least one
foreign protein. For purposes of this specificatIon,
"foreign protein" is intended to mean a protein with
a primary structure as defined by the codons present

15 in the deoxyribonucleic acid sequence which directs
intracellular production of the corresponding amino
acid sequence, and which may or may not include posttranslational modifications. It is further intended
that the term "foreign protein" refers to either the

20 form of the protein as it would be excreted from a
cell or as it may be present in the cell from which
it was not excreted.

While the precise mechanism of regulation is not certain, the promoter/regulator is capable of directing intracellular production of hemoglobin

- and/or other operatively fused gene products upon a drop in oxygen available to the host cell. Also, expression of oxygen-binding proteins, particularly in oxygen-poor environments, can improve
- intracellular production of other proteins.

 The activity of the <u>Vitreoscilla</u> hemoglobin promoter/regulator may be regulated other than by environmental oxygen concentration. During the exponential growth phase of a host cell culture, the

35 addition of a complex nitrogen source, such as yeast

extract, to the cultur allows suppression of the Vitreoscilla hem globin promoter/regulator independently of environmental oxygen concentration. Since the cAMP-CAP complex is important to the activity of the Vitreoscilla hemoglobin promoter/regulator, manipulation of the activity of the crp gene in a host cell is another method of regulation.

In a particularly preferred embodiment, the

10 promoter/regulator contains transcription and
translation initiation and control sequences
substantially equivalent to those for directing
intracellular production of a hemoglobin protein
biologically equivalent to that previously isolated

15 from the filamentous bacterium, Vitreoscilla. By
"substantially equivalent", as used herein, is meant
that a promoter/regulator operates to express a
downstream gene product upon reduction of the level
of oxygen available to the host cell below some

20 critical value.

It is of course intended that the promoter/
regulators of the present invention may control and
initiate transcription and translation of an
unlimited number of endogenous and/or exogenous
25 foreign proteins.

A first preferred portable DNA sequence for the promoter/regulators of the present invention contains at least a portion of the following nucleotide sequence, which reads 5' to 3' and includes the translation initiation sequence <u>ATG</u> (underlined) and some of the nucleotide sequence of the <u>Vitreoscilla</u> structural gene (also underlined):

Hin:

	AAGCTTAACG	GACCCTGGGG	TTAAAAGTAT	TTGAGTTTTG	
	ATGTGGATTA	AGTTTTAAGA			60
	GGCAATAAAG	ATTATAATAA	GTGCTGCTAC	ACCATACTGA	
5	TGTATGGCAA	AACCATAATA			120
	ATGAACTTAA	GGAAGACCCT	CATGTTAGAC	CAGCAAACCA	
	TTAACATCAT	CAAAGCCACT			180
	-		TGGCGTTACC	<u>ATTACCACGA</u>	
	CTTTTTATAA	AAACTTGTTT			240
10	GCCAAACACC		TCCTTTGTTT	GATATGGGTC	200
	GCCAAGAATC	TTTGGAGCAG			300
		TGGCGATGAC TTTGCCAGCT	GGTATTGGCG	GCAGCGCAAA	360
	ACATIGAMAA	TITGCCAGC1			300
15	ATTTTGCCTG AAGCAGGCGR		AATTGCAGTC	<u>AAACATTGTC</u>	420
				•	
		TTGTCGGTCA GGGCGATGCC		GGTGCGATTA	480
		ACATTTTGGA AGATGTGTTT		AAGGCTTATG	540
••	100011000	>> 60> 6> mmm		000000000000000000000000000000000000000	
20		GCCGCTTTCA	GTACGCTCAA	<u>GCGGTTGAA</u> T	600
	00101m111	3300030030	3300maamam	THE THE SAME OF TH	
		CAGCAGCAGT		TTTTACGTCT	660
	###CCC####	CCCS & A A COMM		TTGCCCTGTG	
25		CCGTTGCTGC		1160001616	720
23			•		, 20

GACGTCTTCA GGTGTGCCTT GGCAT

745

Th nucleotide bases represented by the above abbreviations are as follows: A = Adenine, G = Guanine, C = Cytosine, and T = Thymine.

5 The above sequence exhibits homology with certain sequences which are highly conserved in a variety of promoter/regulators. Using conventional numbering, with the underlining showing the homology in the above sequence to the consensus sequence, the -120 consensus sequence or Pribnow box sequence is TATAAT(A/G). The -35 consensus sequence is TTGACA, and the consensus Shine-Dalgarno sequence is AGGAGGTXXX(XX)ATG.

In a preferred embodiment, the above sequence is
operatively fused with at least a portion of a
downstream sequence of nucleotides which code for at
least a portion of the <u>Vitreoscilla</u> hemoglobin
protein which contains at least a portion of the
following amino acid sequence:

20 5 10 Met-Leu-Asp-Gln-Gln-Thr-Ile-Asn-Ile-Ile-

1S 20

Lys-Ala-Thr-Val-Pro-Val-Leu-Lys-Glu-His-

25 30

25 Gly-Val-Thr-Ile-Thr-Thr-Phe-Tyr-Lys-

35

40

Asn-Leu-Phe-Ala-Lys-His-Pro-Glu-Val-Arg-

45 50

Pro-Leu-Phe-Asp-Met-Gly-Arg-Gln-Glu-Ser-

55 60 Leu-Glu-Gln-Pro-Lys-Ala-Leu-Ala-Met-Thr-

65 70

Val-Leu-Ala-Ala-Ala-Gln-Asn-Ile-Glu-Asn-

5 75 80

Leu-Pro-Ala-Ile-Leu-Pro-Ala-Val-Lys-Lys-

90 85 Ile-Ala-Val-Lys-His-Cys-Gln-Ala-Gly-Val-

95 100

10 Ala-Ala-Ala-His-Tyr-Pro-Ile-Val-Gly-Gln-

105 110

Glu-Leu-Leu-Gly-Ala-Ile-Lys-Glu-Val-Leu-

115 120

Gly-Asp-Ala-Ala-Thr-Asp-Asp-Ile-Leu-Asp-

15 125 130

Ala-Trp-Gly-Lys-Ala-Tyr-Gly-Val-Ile-Ala-

135 140

Asp-Val-Phe-Ile-Gln-Val-Glu-Ala-Asp-Leu-

150 145

20 Tyr-Ala-Gln-Ala-Val-Glu

This amino acid sequence is disclosed in Wakabayashi et al., supra, Nature 322:483, 1986. It is presently believed that the protein purified and prepared through the practice of this invention will exhibit a 25 homology of over 80% with this sequence. The protein of this invention has been observed to enhance

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> functioning of a cell in low oxygen envir nments (Khosla and Bailey, unpublished results).

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The amino acids represented by the foregoing abbreviations are as follows:

5	Amino Acid	3-Letter Symbol
	Glycine	Gly
	Alanine	Ala
	Valine	Val
	Leucine	Leu
10	Isoleucine	Ile
	Arginine	Arg
	Lysine	Lys
	Glutamic acid	Glu
	Aspartic acid	Asp
15	Glutamine	Gln
	Asparagine	Asn
	Threonine	Thr
	Serine	Ser
	Cysteine	Cys
20	Methionine	Met
	Phenylalanine	Phe
	Tyrosine	Tyr
	Tryptophan	Trp
	Proline	Pro
25	Histidine	His

It must be borne in mind in the practice of the present invention that the alteration of some amino acids in a protein sequence may not affect the fundamental properties of the protein. Therefore, it 30 is also contemplated that other portable DNA sequences, both those capable of directing intracellular production of identical amino acid sequences and those capable of directing intracellular production of analogous amino acid 35 sequences which also possess oxygen-binding activity, are included within the ambit of the present invention.

It must also be borne in mind in the practice of the present invention that the alteration of some

nucleotide bases in a DNA sequence may not affect the fundamental properties of the coding sequence.

Therefore, it is also contemplated that other analogous portable DNA promoter/regulator sequences

which are operable through changes in oxygen level are included within the ambit of the present invention.

It is contemplated that some of these analogous amino acid sequences will be substantially homologous to native <u>Vitreoscilla</u> hemoglobin while other amino acid sequences, capable of functioning as oxygen-binding proteins, will not exhibit substantial homology to native <u>Vitreoscilla</u> hemoglobin. By "substantial homology" as used herein, is meant a degree of homology to native <u>Vitreoscilla</u> hemoglobin in excess of 50%, preferably in excess of 80%.

Similarly, it is contemplated that some of these analogous DNA sequences will be substantially homologous to the sequence set forth above, while other DNA sequences, capable of functioning as the promoter/regulator described above, will not exhibit substantial homology to the sequence outlined above.

As noted above, the portable DNA sequences of the present invention may be synthetically created, by

25 hand or with automated apparatus. It is believed that the means for synthetic creation of these polynucleotide sequences are generally known to one of ordinary skill in the art, particularly in light of the teachings contained herein. As examples of the current state of the art relating to polynucleotide synthesis, one is directed to Maniatis et al.,

Molecular Cloning—A Laboratory Manual, Cold Spring Harbor Laboratory (1984), and Horvath et al., An Automated DNA Synthesizer Employing Deoxynucleoside

3'-Phosphoramidit s, Methods in Enzymology 154:313-326, 1987, hereby incorporated by reference.

Additionally, the portable DNA sequence may be a fragment of a natural sequence, i.e., a fragment of a polynucleotide which occurred in nature and which has been cloned and expressed for the first time by the present inventors. In one embodiment, the portable DNA sequence is a restriction fragment isolated from a genomic library. In this preferred embodiment, the genomic library is created from the bacterium Vitreoscilla. In other alternative embodiments, the portable DNA sequence is isolated from other genomic and cDNA libraries.

While it is envisioned that the portable DNA 15 sequences of this invention may desirably be inserted directly into the host chromosome, the present invention also provides a series of vectors, each containing at least one of the portable DNA sequences described herein. It is contemplated that additional 20 copies of the portable DNA sequence may be included in a single vector to increase a host cell's ability to produce large quantities of the desired oxygenbinding protein. It is also envisioned that other desirable DNA sequences may also be included in the 25 vectors of this invention. Further, the invention may be practiced through the use of multiple vectors, with additional copies of at least one of the portable DNA sequences of this invention and perhaps other desirable DNA sequences.

30 In addition, the cloning vectors within the scope of the present invention may contain supplemental nucleotide sequences preceding or subsequent to the portable promoter/regulator and/or DNA sequence.

These supplemental sequences are those that will not

adversely interfer with transcription of the portable promoter/regulator and/or any fused DNA sequence and will, in some instances, enhance transcription, translation, posttranslational processing, or the ability of the primary amino acid structure of the resultant gene product to assume an active form.

A preferred vector of the present invention is set forth in Figure 1. This vector, pUC19/pRED2,

10 contains the preferred nucleotide sequence which codes for the amino acids set forth above. Vector pUC19/pRED2 cells are on deposit in the American Type Culture Collection ("ATCC") in Rockville, Maryland under Accession No. 67536.

- 15 A preferred nucleotide sequence encoding the

 Vitreoscilla hemoglobin protein and adjacent

 Vitreoscilla sequences described above is identified in Figure 1 as region A. The above nucleotide sequence reads counterclockwise through region A of
- 20 Figure 1. Plasmid pUC19/pRED2 may also contain supplemental nucleotide sequences preceding and subsequent to the preferred DNA sequence in region A, such as terminators, enhancers, attenuators and the like. For proteins to be exported from the
- 25 intracellular space, at least one leader sequence and any other DNA sequences necessary or preferred for appropriate transcription and subsequent translation of the vector DNA may be included within the scope of this invention.
- 30 In a preferred embodiment, cloning vectors containing and capable of expressing the portable DNA sequence of the present invention contain various operational elements in addition to or instead of the promoter/regulator disclosed and claimed herein.

These "operational elements" may include at least one promoter, at least one sequence that acts as expression regulator, and at least one terminator codon, at least one leader sequence, and any other

5 DNA sequences necessary or preferred for appropriate transcription and subsequent translation of the vector DNA.

Additional embodiments of the present invention are envisioned as employing other known or currently 10 undiscovered vectors which would contain one or more of the portable DNA sequences described herein. particular, it is preferred that these vectors have some or all of the following characteristics: (1) possess a minimal number of host-organism 15 sequences; (2) be stable in the desired host; (3) be capable of being present in a high copy number in the desired host; (4) possess a regulatable promoter; and (5) have at least one DNA sequence coding for a selectable trait present on a portion of the plasmid 20 separate from that where the portable DNA sequence will be inserted. Alteration of vectors to meet the above criteria are easily performed by those of ordinary skill in the art in light of the available literature and the teachings herein. It is to be 25 understood that additional cloning vectors may now exist or will be discovered which have the aboveidentified properties and are therefore suitable for use in the present invention and these vectors are also contemplated as being within the scope of this 30 invention.

As set forth in Example 1, an <u>E. coli</u> vector system is a preferred embodiment. Various cloning vehicles are required for the range of host cells and organisms suitable for insertion of the portable DNA sequences of the present invention, as set forth

below. In light f the available literature, choice of such a cloning vehicle, if necessary, is within the ordinary skill in the art.

- Additional bacterial hosts are suitable, including,

 5 without limitation: bacteria such as members of the
 genera <u>Bacillus</u>, <u>Pseudomonas</u>, <u>Alcaligenes</u>,

 <u>Streptococcus</u>, <u>Lactobacillus</u>, <u>Methylophilus</u>,

 <u>Xanthomonas</u>, <u>Corynebacterium</u>, <u>Brevibacterium</u>,

 <u>Acetobacter</u>, and <u>Streptomyces</u>.
- 10 Examples of suitable eucaryotic host microorganisms would include fungi, yeasts such as <u>Saccharomyces</u> and <u>Candida</u>, and molds such as <u>Asnergillus</u>, <u>Pennicillim</u> and <u>Cephalosporium</u>.
- It is envisioned that the scope of this invention

 15 would cover expression systems in eucaryotic

 microorganisms and host cultured cells derived from

 multicellular organisms, including animals, insects

 and plants, which are grown in the presence of

 oxygen. The promoter/regulator of the present
- invention is especially useful in a host which switches from low to very high expression activity upon reduction of dissolved oxygen concentration in the medium. Such expression systems need not be derived from Vitreoscilla.
- 25 Various vector systems will be suitable for these and other desirable hosts, including plasmids, viruses and bacteriophages. The following, noninclusive list of cloning vectors is believed to set forth vectors which can easily be altered to meet the above
- 30 criteria and are therefore preferred for use in the present invention. Such alterations are easily performed by those of ordinary skill in the art in

light of the availabl literature and the teaching herein.

For example, many selectable cloning vectors have been characterized for use in <u>E. coli</u>, including

5 pUC8, pUC9, pBR322, pGW7, placIq, and pDP8, Maniatis et al., supra. A bifunctional vector that replicates in <u>E. coli</u> and can also be used in <u>Streptomyces</u> is pKC462a. Suitable vectors for use in <u>Bacillus</u> include: pUB110, pSA0501, pSA2100, pBD6, pBD8, and pT127, Ganesan and Hock, eds., <u>Genetics and Biotechnology of Bacilli</u>, Academic Press 1984. In <u>Pseudomonas</u>, RSF1010, Pms149, pKT209, and RK2 are suitable; some of these vectors are useful in a wide

15 Agrobacterium and Xanthomonas. For <u>Saccharomyces</u>, it is possible to use YEp24, Ylp5, and YRp17, Botstein and Davis, <u>Molecular Biology of the Yeast</u>

<u>Saccharomyces</u> (Strathern <u>et al</u>., eds.), Cold Spring Harbor Laboratory, 1982. In mammalian systems

range of gram-negative bacteria including

20 retrovirus vectors such as those derived from SV40 are typically used.

Synthesis and/or isolation of necessary and desired component parts of cloning vectors, and their assembly is believed to be within the duties and tasks performed by those with ordinary skill in the art and, as such, are capable of being performed without undue experimentation.

In construction of the cloning vectors of the present invention, it should additionally be noted that

30 multiple copies of the promoter/regulator with any fused gene sequences and/or of the portable DNA sequence coding for the oxygen-binding protein and its attendant operational elements as necessary may be inserted into each vector. In such an embodiment,

the host organism would produce greater amounts per vector of the desired cloned protein. The number of multiple copies of the DNA sequence which may be inserted into the vector is limited only by the ability of the resultant vector, due to its size, to be transferred into and replicated and expressed in an appropriate host.

Additionally, it is preferred that the cloning vector contain a selectable marker, such as a drug 10 resistance marker or other marker which causes expression of a selectable trait by the host. In a particularly preferred embodiment of the present invention, the gene for ampicillin resistance is included in vector pUC19/pRED2. Such a drug 15 resistance or other selectable marker is intended in part to facilitate in the selection of transformants. Additionally, the presence of such a selectable marker on the cloning vector may be of use in keeping contaminating microorganisms from multiplying in the 20 culture medium. In this embodiment, such a pure culture of the transformed host organisms would be obtained by culturing the organisms under conditions which require the induced phenotype for survival.

It is noted that the portable DNA sequence of the
present invention may themselves be used as a
selectable marker, in that they provide enhanced
growth characteristics in low oxygen circumstances,
and also engender an easily visible reddish tint in
the host cells.

30 The promoter/regulators of this invention are capable of controlling expression of proteins or, thereby, of controlling synthesis of metabolites normally made by a cell, or of natural or unnatural metabolites and proteins expressed in a cell via genetic

manipulation. This would include heterol g us proteins—either intracellular or extracellular—as well as biopolymers such as polysaccharide materials, simpler metabolites such as amino acids and nucleotides, antibiotics and other chemicals produced by living cells or cellular biocatalysts.

The oxygen-binding proteins of the present invention, prepared by the recombinant-DNA methods set forth herein, will enable increased research into the 10 growth of organisms in oxygen-poor environments. In addition, the oxygen-binding proteins of the present invention are useful in enhancing oxygen supply to cells or in other oxygen-utilizing processes (Adlercreutz et al., Biocatalyst in Organic 15 Synthesis, Symposium of the Working Party on Immobilized Biocatalysts of the European Federation of Biotechnology, Abstracts, p.18, 1985), and for binding and separating oxygen from other fluids or gases (Bonaventura et al., Underwater Life Support 20 Based on Immobilized Oxygen Carriers, Applied Biochemistry and Biotechnology 9: 65-80, 1984). Furthermore, the oxygen-binding proteins of this invention are capable of increasing production of cells, or of proteins or metabolites normally made by 25 a cell, or of natural or unnatural metabolites and proteins expressed in a cell via genetic manipulation. This would, as described above, include heterologous proteins, biopolymers, simpler metabolites, antibiotics, and other chemicals 30 produced by living cells or cellular biocatalysts.

The protein products of this invention also have applications as diverse as enhancing certain oxygen-requiring steps in fermentation, enzymatic degradation, toxic chemical waste treatment, brewing

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and particular oxidative reactions and transformations such as steroid conversions.

This invention also relates to a recombinant-DNA method for the production of oxygen-binding proteins.

5 Generally, this method includes:

- (a) preparing a portable DNA sequence capable of directing a host cell or microorganism to produce a protein having oxygen-binding activity;
- (b) transferring the portable DNA sequence directly into the host, or cloning the portable DNA sequence into a vector capable of being transferred into and replicating in a host cell or microorganism, such vector containing operational elements for the portable DNA sequence;
- (c) transferring the vector containing the portable DNA sequence and operational elements into a host cell or microorganism capable of expressing the oxygen-binding protein;
- (d) culturing the host microorganism under conditions appropriate for replication and propagation of the vector and/or expression of the protein; and
- (e) in either order:
 - (i) harvesting protein; and(ii) permitting the protein to assumean active structure whereby itpossesses oxygen-binding activity.

In this method, the portable DNA sequences are those synthetic or naturally-occurring polynucleotides described above. In a preferred embodiment, the

portable DNA sequence codes for at least a portion of the <u>Vitreoscilla</u> hemoglobin protein described above.

This invention also relates to a recombinant-DNA m thod for the use of thes promoter/regulators.

5 Generally, this method provides a process for subjecting the expression of a selected DNA sequence to external control under given environmental conditions which comprises the steps of:

- (a) providing at least one selected isolated

 10 structural gene that is transcriptionally and/or
 translationally responsive to a <u>Vitreoscilla</u>
 hemoglobin promoter/regulator DNA sequence under the
 given environmental conditions; and
- (b) operatively fusing the selected structural 15 gene with said promoter/regulator DNA sequence.

It is envisioned that the portable DNA sequences may be inserted directly into the host chromosome, or alternatively may utilize a vector cloning system. The vectors contemplated as being useful in the present method are those described above. In a preferred embodiment, the cloning vector pUC19/pRED2 is used in the disclosed method.

A vector thus obtained may then be transferred into the appropriate host cell or organism. It is

25 believed that any microorganism having the ability to take up exogenous DNA and express those genes and attendant operational elements may be chosen.

Particular hosts which may be preferable for use in this invention include those described above.

30 Methods for transfer of vectors into hosts are within the ordinary skill in the art. For ultimate expression in certain microorganisms such as yeast, it may be desirable that the cloning vector be first transferred into another microorganism such as

Escherichia coli, where the vector would be allowed t replicate and, from which the vector would be obtained and purified after amplification, and then transferred into the yeast for ultimate expression of the oxygen-binding protein.

The host cells or microorganisms are cultured under conditions appropriate for the expression of the oxygen-binding protein. These conditions are generally specific for the host organism, and are readily determined by one of ordinary skill in the art, in light of the published literature regarding the growth conditions for such organisms.

In one embodiment, conditions necessary for the regulation of the expression of the DNA sequence, 15 dependent upon any operational elements inserted into or present in the vector, would be in effect at the transformation and culturing stages. The cells are grown to a high density in the presence of appropriate regulatory conditions which inhibit the 20 expression of the DNA sequence. When optimal cell density is approached, the environmental conditions are altered to those appropriate for the expression of the portable DNA sequence. It is thus contemplated that the production of a cloned protein 25 will occur in a time span subsequent to the growth of the host cells to near optimal density, and that the resultant cloned protein product would be harvested, if desired, at some time after the regulatory conditions necessary for its expression were induced.

30 Where the operational elements used are in the promoter/regulator sequence of this invention, these conditions are as follows. The cells are grown to a high density in the presence of appropriate levels of oxygen which inhibit the expression of the DNA

sequence. When optimal cell density is approached, the environmental oxygen level is altered to a lower value appropriate for the expression of the portable DNA sequence. Levels from less than in a 1% oxygensaturated solution to oxygen saturated are within the scope of this invention. It is thus contemplated that the production of any desired fused product will occur in a time span subsequent to the growth of the host cells to near optimal density, and that the resultant product would be harvested, if desired, at some time after the oxygen level necessary for its expression were reached.

If harvesting of the oxygen-binding protein products of the present invention is desired, it may be done prior or subsequent to purification and prior or subsequent to assumption of an active structure.

It is currently believed that some percentage of the oxygen-binding proteins of the present invention will assume their proper, active structure upon expression 20 in the host cell or organism. If desired, the oxygen-binding protein may be transported across a cell membrane. This will generally occur if DNA coding for an appropriate leader sequence has been linked to the DNA coding for the recombinant protein. 25 The structures of numerous signal peptides have been published. It is envisioned that these leader sequences, included in or added to at least some portion of the portable DNA as necessary, will direct intracellular production of a fusion protein which 30 will be transported through the cell membrane and will have the leader sequence cleaved upon release from the cell.

Additional uses of the oxygen-binding proteins of the present invention are envisioned. The purified

proteins and/or the wh le cells and/or extracts of the cells of the present invention themselves may be used to bind to oxygen or proteins and thus could function somewhat as erythrocytes.

The present invention may also be used as a method for transporting and enhancing oxygen supply to cells or in other oxygen-utilizing processes by delivering the oxygen-binding proteins -- isolated in lysates and crude cell preparations, purified from extracts, in synthetic sequences, or in whole cells containing the proteins -- where desired. It is envisioned that the protein products of the present invention could valuably be added to media for culturing cells and thereby enhance the transport of oxygen.

It is also envisioned that the proteins of the present invention may be used for binding and separating of oxygen from fluids such as seawater and from other gases.

20 It is understood that application of the teachings of the present invention to a specific problem or environment will be within the capabilities of one having ordinary skill in the art in light of teachings contained herein. Examples of the products of the present invention and representative processes for their isolation, use and manufacture appear below.

INDUSTRIAL APPLICABILITY

The products and processes of the present invention

find usefulness in a range of medical, laboratory and
industrial applications. The invention provides
metabolically engineered cells with enhanced growth
characteristics for increasing production of various

proteins or metabolites by those cells. The invention further provides a method for subjecting expression of a certain DNA sequence to external control under given environmental conditions. Also provided are recombinant-DNA fusion gene products, expression vectors, and nucleotide base sequences for the practice of the invention. The products and processes of the present invention find applications in a range of aerobic processes, such as manufacture of cloned proteins and synthesis of metabolites, chemical production by fermentation, enzymatic degradation, waste treatment, brewing and a range of oxidative reactions.

EXAMPLES

15 EXAMPLE 1 - CLONING AND EXPRESSION OF HEMOGLOBIN FROM VITREOSCILLA IN ESCHERICHIA COLI.

Materials and Methods. Vitreoscilla sp. (Murray strain no. 389) was obtained from Dr. Webster (Department of Biology, Illinois institute of Technology, Chicago, Illinois 60616, USA), and grown in a medium containing 1.5% yeast extract, 1.5% peptone, and 0.02% sodium acetate (pH 8.0 with NaOH).

E. coli JM101 were obtained from the laboratory of Dr. Simon (Division of Biology, California 25 Institute of Technology, Pasadena, California 91125, USA), and grown in L broth containing 1% Bactotryptone, 0.5% yeast extract and 1% sodium chloride.

Plasmid pUC19 (Yanisch-Perron et al., Improved

30 M13 phase cloning vectors and host strains:

nucleotide sequences of m13mp18 and nUC19 vectors,

Gene 33: 103-109, 1985) packaging kits were purchased from Pharmacia. All restriction enzymes, T4

polynucleotide kinase and T4 ligase were from New England Biolabs or Bethesda Research Laboratories.
Calf intestine alkaline ph sphatase was from Pharmacia. Mixed oligonucleotide probes were synthesized with an Applied Biosystems synthesizer. Kodak XAR5 x-ray film was used for autoradiography. Geneclean kits were purchased from Biolol. All other chemicals were of analytical grade.

Vitreoscilla genomic DNA was isolated according to

the protocol of Silhavy et al., Experiments with gene
fusions, Cold Spring Harbor Laboratory (1984),
specifically incorporated herein. HindIII-digested
Vitreoscilla DNA was ligated into the phosphatased
HindIII site of pUC19 and transformed into JM101.

- 15 Recombinant colonies and plaques were transferred on nitrocellulose filters as described in Maniatis, et al., Molecular cloning--a laboratory manual, Cold Spring Harbor Laboratory (1982) and specifically incorporated herein. Rapid plasmid isolation from
- 20 recombinant colonies were done according to Silhavy et al, supra. Digested fragments of plasmid DNA or fractions of genomic DNA were isolated from agarose gels using Geneclean kits. E. coli cells were transformed by the CaCl2 method of Silhavy et al.,
- 25 <u>supra</u>. Plasmid uptake was induced by heat-shocking chilled competent cells at 37°C for 5 minutes.

For Southern hybridizations the reagents suggested in Dupont catalog No. NEF-976, <u>Protocols for electrophoretic and capillary transfer of DNA and RNA</u>

- DNA and RNA hybridization and DNA and RNA
 rehybridization (1985), specifically incorporated
 herein, were used, whereas for colony and plaque
 hybridizations those described in Maniatis et al.,
 supra, were used. Filters were prehybridized at 45-
- 35 50 C for 2-4 hours and hybridized at 30°C for 20-24

hours. 200 picomoles oligonucleotide kinased with 200 microCi (32_p)ATP (sp. act. 7000 Ci/mmol) were used as probe. Filters were washed in 2 X SSC, 0.1% SDS at room temperature (3 X 5 minutes) and at 46°C (for the C-terminal probe) and 50°C (for the N-terminal probe) prior to autoradiography.

SDS-polyacrylamide gel electrophoresis was done according to standard protocols, Laemmli, Cleavage of structural proteins during the assembly of the head of bacteriophage T4, Nature 227:680-685, 1970, specifically incorporated herein, with a 12.5% resolving gel. Protein in the gel was visualized by the silver staining method of Merril et al., Ultrasensitive stain for proteins in polyacrylamide gels shows regional variation in cerebrospinal fluid proteins, Science 211:1437-1438, 1983.

Results. Three sets of mixed oligonucleotide probes were synthesized which had a predicted homology to two domains in the hemoglobin gene, one N-terminal 20 and one C-terminal. A pUC19-HindIII library of Vitreoscilla DNA was test-plated on rich plates with ampicillin, X-gal and IPTG. More than 70% of the colonies were probable recombinants, as estimated by visual inspection. About 10,000 colonies were then 25 screened. Three positives were identified. Because of the high density of colonies on the plate, these, along with their immediate clones from each group, were assayed by rapid isolation and HindIII digestion. One of these, pRED1, had three inserted 30 fragments including a 2.2kb one. Subsequent digestion of this plasmid with various endonucleases and Southern hybridization of the resulting DNA bands did confirm that the 2.2kb band did indeed contain the entire hemoglobin gene, since no HindIII sites

are expected to exist upstream or downstream of the regions spanned by the oligomeric probes.

The HindIII fragment from pRED1 that contained th hemoglobin structural gene was purifi d and

5 reinserted by standard protocols into pUC19 in both orientations (pRED2 and pRED3). E. coli cells containing plasmids pRED1, pRED2, pRED3 and pUC19 as well as Vitreoscilla cells were grown to stationary phase and cell extracts were assayed on an

10 SDS-polyacrylamide gel for the existence of the hemoglobin polypeptide. The hemoglobin was expressed as a major cellular protein in all recombinant cells. Since both plasmids pRED2 and pRED3 express about equal amounts of this polypeptide, it is presently believed that the gene is probably expressed from its natural promoter in E. coli.

The restriction map of plasmid pRED2 is shown in Figure 1.

EXAMPLE 2 - COMPLETE NUCLEOTIDE SEQUENCE.

20 Determination of the sequence of the relevant region of the fragment isolated from the <u>Vitreoscilla</u> genomic library was accomplished as follows:

The HindIII-SphI fragment from plasmid pRED2 which contains the structural gene and adjacent sequences

25 was subcloned into pUC19 (purchased from Bethesda Research Labs) to obtain plasmid pRED4. An MluI site was identified, by restriction mapping the resulting plasmid, which breaks up the HindIII-SphI insert into two fragments which were individually sequenced using conventional protocols (Maxam and Gilbert, Sequencing end-labeled DNA with base-specific chemical cleavages, Methods in Enzymology 65: 499-560, 1980; Iverson and Dervan, Adenine specific DNA chemical

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sequencing reaction, Nuclear Acids Research 15:
7823-7830, 1987).

The nucleotide sequence of the important portion of the HindIII-SphI fragment is as listed below. It

5 includes a putative <u>E. coli</u> promoter, ribosome binding site, the complete VHb structural gene (start and stop codons are underlined) and a putative <u>E. coli</u> transcription terminator (Khosla and Bailey, <u>The Vitreoscilla hemoglobin gene: molecular cloning</u>

10 nucleotide sequence and genetic expression in <u>Escherichia coli</u>, Mol. & Gen. Genet., in press).

AAGCTTAACG GACGCTGGGG TTAAAAGTAT TTGAGTTTTG ATGTGGATTA AGTTTTAAGA 60 GGCAATAAAG ATTATAATAA GTGCTGCTAC ACCATACTGA 15 TGTATGGCAA AACCATAATA 120 ATGAACTTAA GGAAGACCCT CATGTTAGAC CAGCAAACCA TTAACATCAT CAAAGCCACT 180 GTTCCTGTAT TGAAGGAGCA TGGCGTTACC ATTACCACGA CTTTTTATAA AAACTTGTTT 240 20 GCCAAACACC CTGAAGTACG TCCTTTGTTT GATATGGGTC GCCAAGAATC TTTGGAGCAG 300 CCTAAGGCTT TGGCGATGAC GGTATTGGCG GCAGCGCAAA ACATTGAAAA TTTGCCAGCT 360 ATTTTGCCTG CGGTCAAAAA AATTGCAGTC AAAOGATTGTC 25 AAGCAGGCGR GGCAGCAGCG 420 CATTATCCGA TTGTCGGTCA AGAATTGTTG GGTGCGATTA

AAGAAGTATT GGGCGATGCC

G	CAACCGATG	ACATTTTGGA	CGCGTGGGGC	AAGGCTTATG	
G	CGTGATTGC	AGATGTGTTT			540
A	TTCAAGTGG	AAGCAGATTT	GTACGCTCAA	GCGGTTGAA <u>T</u>	
A	<u>A</u> AGTTTCAG	GCCGCTTTCA			600
5 G	GACATAAAA	AACGCACCAT	AAGGTGGTCT	TTTTACGTCT	
G	SATATTTACA	CAGCAGCAGT			660
1	TGGCTGTTG	GCCAAAACTT	GGGACAAATA	TTGCCCTGTG	
1	PAAGAGCCCG	CCGTTGCTGC			720
G	SACGTCTTCA	GGTGTGCCTT	GCCAT		745

10 EXAMPLE 3 - GROWTH ENHANCEMENT IN E. COLI WITH pRED2: SHAKE FLASK CULTURES.

In this Example, the growth behavior of <u>E. coli</u> cells producing active <u>Vitreoscilla</u> hemoglobin was compared to that of control strains grown under identical conditions. The following strains were studied:

(1) JM101:pRED2; (2) JM101:pUC9; and (3) JM101.

Plasmids pUC9 and pUC19 are essentially identical except for a difference in one restriction site unrelated to the insert or to any of the functional properties of the plasmid.

Experimental protocol: Cells were grown at 37°C in a complex medium containing 1% (W/V) bactotryptone, 0.5% (W/V) yeast extract, 0.5% (W/V) NaCl, 0.3% (W/V) K2HPO4 and 0.1% (W/V) KH2PO4 (pH 7.0). Plasmidcontaining cells were grown in the presence of 100 mg/L ampicillin. In each case the shake-flask was inoculated with a 1% (V/V) dose of concentrated nutrient broth containing 430 g/L glucose, 5 g/L yeast extract, 110 g/L (NH4)2SO4, 8 g/L MgSO4 · 7H20, 0.27 g/L FeCl3 · 6H20, 0.02 g/L ZnCl2 · 4H2O, 0.02 g/L CaCl2 · 2H2O, 0.02 g/L Na2MOO4 · 2H2O, 0.01 g/L

 $CusO_4 \cdot 5H_2O$, 0.005 g/L H_3BO_3 , 0.1% (V/V) conc. HC1, 4.2 mg/L riboflavin, 54 mg/L pantothenic acid, 60 mg/L folic acid. This formulation has been successfully used on a previous occasion to grow 5 stationary cells to a high density in a fedbatch mode. The cells were then allowed to grow further until stationary phase was reached again. Optical density was measured at 600 nm on a Bausch & Lomb Spectronic 21 spectrophotometer. Dry weights were 10 measured by spinning 10 mL samples at 4°C, washing once with distilled water and subsequently drying the resuspended sample at 100°C overnight. The heme content of the cells was assayed according to the method of Lamba & Webster (Lamba & Webster, Effect of 15 growth conditions on yield and heme content of Vitreoscilla, Journal of Bacteriology 142: 169-173, 1980), and the hemoglobin activity was measured by the method of Webster & Liu (Webster and Liu, Reduced nicotinamide adenine dinucleotide cytochrome o 20 reductase associated with cytochromic o purified from Vitreoscilla, Journal of Biological Chemistry, 249: 4257-4260, 1974).

Results. The growth properties, heme content and hemoglobin activity of the three strains aredocumented in the Table below.

		<u>JM1</u>	01:pRED2 J	M101:pUC9	<u>JM101</u>
	1.	OD ₆₀₀ before nutrient replenishment	0.937	0.737	0.945
		2 Cp 20112 Dilliacite	0.35.	0	0.745
5	2.	OD ₆₀₀	1.230	0.880	0.985
	3.	<pre>max. attained dry wt.</pre>	1.5g/L	0.85g/L	lg/L
	4.	relative heme content	5.5	1	**
10	5.	relative hemoglobi activity	n 5	1	**
	6.	specific growth rate*	0.04/h	0.01/h 0	0.009/h

*mean value following additional feeding of shake15 flasks as described above

EXAMPLE 4

GROWTH ENHANCEMENT OF E. COLI WITH DRED2. A typical high-cell density fermentation is of a fed-20 batch type. The optimal rate of nutrient addition, and consequently the productivity, is ultimately limited by the rate at which cells can aerobically catabolize the carbon source without generating growth-inhibitory metabolites such as acetate and 25 lactate (Zabriskie and Arcuri, Factors influencing productivity of fermentations employing recombinant microorganisms, Enzyme and Microbial Technology 8: 706-717, 1986; Tsai et al, The effect of organic nitrogen and glucose on the productivity of 30 recombinant insulin-like growth factor in high cell density Escherichia coli fermentations, Journal of Industrial Microbiology 2: 181-187, 1987). In this experiment, we compare the growth properties of the recombinant strain (JM101:pRED2) expressing 35 <u>Vitreoscilla</u> hemoglobin with similar plasmidcontaining (JM101:pUC9) and plasmid-free (JM101)

^{**}not assayed

strains under typical fed-batch fermentation conditions.

Materials and Methods:

Cells were grown in a New Brunswick Microferm 5 fermentor at 37+0.5°C and a pH of 7±0.05 with an initial working volume of 2.5 L. A constant air-flow rate of 4.5 L/min and agitator speed of 300 rpm were maintained throughout each run. Silicone anti foam AF60 was used to control foaming. The batch medium 10 and feed medium 1 listed in Table 2 in Tsai et al., supra. were used. Growth following inoculation was in batch mode. After batch stationary phase was reached, continuous feeding was initiated using feed medium 1 at a flow rate of 10 mL/hr. For plasmid-15 containing cells, 100 mg/L ampicillin was used. all cases, the dissolved oxygen (DO) levels remained fairly constant around 5% of air saturation for most of the run except during the early log phase and towards batch stationary phase.

20 Results:

The growth parameters measured for the three strains are listed below. Batch stationary phase refers to conditions before continuous feeding was started.

		<u>JM101</u>	JM101:pUC9	JM101:pRED2
25	Batch log-phase growth rate (h-1)	0.95	0.73	0.95
	Batch stationary- phase dry cell mass (g/L)	2.6	1.6	2.6
30	Fed-batch log-phase growth rate (h ⁻¹)	0.056	0.033	0.066
	Final dry cell man	ss 5.8	2.8	5.9

Further, the respiratory behavior of JM101:pRED2 was improved compared to the control strains at low DO levels, as observ d in a Gilson respirometer.

Conclusion:

5 Cells containing <u>Vitreoscilla</u> hemoglobin grow faster and to higher densities than comparable plasmid-containing controls.

EXAMPLE 5

EXPRESSION OF VITREOSCILLA HEMOGLOBIN (VHb) IN E. COLI UNDER THE REGULATION OF OTHER PROMOTERS 10 In Examples 1, 3, and 4 above, the expression of hemoglobin is under the regulation of its native oxygen-regulated promoter. Hence, it is not possible to modulate independently the dissolved oxygen 15 concentration (DO) and the intracellular VHb level. In order to overcome this, the inventors attempted to express this protein under the control of other regulatable promoters which are functional in E. coli, such as trp (Russell and Bennett, 20 Construction and analysis of in vitro activity of E. coli promoter hybrids and promoter mutants that alter the -35 to -10 spacing, Gene 20: 231-243, 1982) and tac (deBoer et al., The tac promoter: a functional hybrid derived from the trp and lac promoters, Proc. 25 Natl. Acad. Sci. USA 80: 21-25, 1983).

Materials and Methods:

Plasmid pRED4 (see Example 2) was linearized with HindIII and treated with exonuclease Bal31 to generate 5'end deletions in the HindIII-SphI VHb 30 fragment (Maniatis et al., supra). After digestion with SphI, the resulting VHb fragments were cloned into HindIII-SphI digested pUC19. The positions of the deleted end-points were identified by sequencing (protocol similar to that in Example 2).

trp and tac promoters and the chloramphenicol acetyl transferase gene (CAT) were purchased from Pharmacia, Inc. Oligonucleotides were synthesized at California Institute of Techn logy using an Applied Biosystems
5 DNA synthesizer. All DNA nzymes were obtained from vendors.

The functional assay for the VHb gene product is as described in Webster and Liu, <u>supra</u>.

Cells were pelleted at 4°C and resuspended in 100 mM

Tris (pH 7.5), 50 mM NaCl. This cell suspension was sonicated at 75 W for 3 min. on ice. After spinning in at 12,000 g for 10 min., the supernatant was collected and assayed for VHb. Total protein content was estimated using the Bradford assay kit from

BioRad Inc. VHb activity is reported as delta-A419436/mg total protein.

Results:

One of the deletions, pRED302, mapped 2 base-pairs upstream of the ATG start codon for the VHb

20 structural gene. This deletion was used for further work. The EcoRI/BamHI trp promoter cartridge was cloned upstream of the truncated VHb fragment. The following ribosome binding site was synthesized:

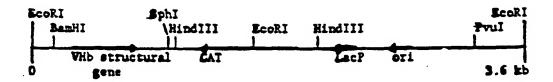
5' GATCCCGGGTCTAGAGGA 3' GGCCCAGATCTCCT

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and inserted between the BamHI and nuclease-blunted XbaI sites to give rise to a trp promoter-controlled VHb expression system. The CAT gene (Alton and Vapnek, Nucleotide sequence analysis of the Chloramphenical resistance transposon Tn9, Nature 282: 864-869, 1979) was inserted downstream and under the control of the lac promoter available on

this pUC19-based plasmid. This gene product can be conveniently assayed (Neumann et al, Novel rapid assay for chloramphenicol acetyltransferase gene expression, BioTechniques 5: 444-447, 1987) and 5 henc serv s as a useful reporter. Finally, the B-lactamase gene on this pUC19-based plasmid was deleted by digestion and religation with PuS. The purpose of this step is to eliminate the presence of a plasmid-encoded periplasmic protein. The plasmid 10 thus obtained was called pHbCAT and was transformed into JM101. As a control, the CAT gene was cloned downstream and under the control of the lac promoter in pUC19. The B-lactamase gene was identically deleted. This plasmid was called pCAT. 15 restriction maps and the anticipated sequence of relevant regions of these two plasmids are shown below.

PHOCAT (3.6 kb)



EcoRi

5' GAATT CCCCT GTTGA CAATT AATCA TCCAA CTAGT TAACT
20 AGTAC

BamHI

GCAGC TTGGC TGCAG GTCGA CGGAT CCCGG GRCTA GAGGA AGTCT

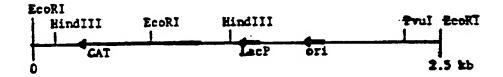
Start codon of VHb

CATGT TAGAC (same as in Example 3 up to SphI site)

The sequence of the region spanning between EcoRI and the start of the VHb structural gene is shown above.

It includes th trp promoter and a synthetic ribosome
binding site.

pCAT (2.5 kb)



The effect of tryptophan (repressor) and indole

5 acrylic acid (gratuitous inducer) on VHb levels in

JM101/pHbCAT are shown in the Table below. In these
experiments, cells were grown to mid-log in minimal
medium containing 3 g/L glycerol, 3 g/L Casamino
acids, and the appropriate amount of indole-acrylic

10 acid or tryptophan.

	Host:	Tryptophan	Indole-acry	ylic Specific Hb
	Plasmid	(mg/L)	acid (mg/	L) Activity*
	JM101:pCAT			3.4×10^{-3}
5	JM101:pHbCA	T 20		6.2 x 10 ⁻³
	n	4		18.3×10^3
	ŧı			31.5 x 10 ⁻³
	•1		1	29.8 x 10 ^{.3}
.10	**		2.53	6.5 x 10 ⁻³
	11		5	47.0 x 10 ⁻³
	n		10	36.6 x 10 ⁻³

Note: *(delta-A₄₁₉₄₃₆/mg total soluble protein)

To express the VHb gene under the control of the tac

15 promoter, an expression plasmid was made using a

HindIII-BamHI tac promoter cartridge, the BamHI/SphI

fragment from pHbCAT, and the HindIII-SphI digested

fragment of the vector pBR322 (Bolivar et al.,

Construction and characterization of new cloning

vehicles. II. A multipurpose cloning system, Gene

2: 95-113, 1977).

With this construct (pINT1), the level of redness of cells correlated well with varying amounts of the gratuitous inducer IPTG, indicating that the gene product synthesis was under the control of tac. The advantages of this expression system are:

- a. Higher expression of VHb, and
- b. Ability to use complex medium for growth.

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EXAMPLE 6

GROWTH OF E. COLI - EXPRESSING VHb UNDER THE REGULATION OF OTHER PROMOTERS

The aim of this experiment was to demonstrate the growth effects of VHb on <u>F. coli</u>. In thes cases, VHb is expressed using promoters different from the native VHb oxygen-regulated promoter. The strains:plasmids used are:

- 1. HB101:pBR322 (pBR322 from BRL)
- 2. JM101:pINT1 (pINT1 discussed in Example 5). The two hosts have nearly identical genotypes, the major difference being the presence of an F' factor in JM101 which harbors the lacIq gene. This gene is necessary to keep a strong promoter like tac under control.

The following media recipes shall be henceforth referred to in the appropriate annotated form:

1X LB: 10 g/L Bactotryptone, 5 g/L Yeast
Extract, 5 g/L NaCl, 3 g/L K₂HPO₄, 1
g/L KH₂PO₄, 100 mg/L Ampicillin

2X LB: 20 g/L Bactotryptone, 10 g/L Yeast Extract, 5 g/L NaC1, 3 g/L K₂HPO₄, 1 g/L KH₂PO₄, 100 mg/L Ampicillin

25 Extract, 5 g/L NaCl, 3 g/L K₂HPO₄, 1 g/L KH₂PO₄, 100 mg/L Ampicillin.

The experiment was conducted as follows. Single colonies of the two strains listed above were inoculated into 5 mL 1X LB in a culture tube and 30 grown overnight at 37°C.

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0.5 mL of th appropriate inoculum was transferred into 250 mL culture flasks c ntaining 50 mL medium as follows:

1) HB101:pBR322 : 2X LB 5 2) HB101:pBR322 : 5X LB 3) JM101:pINT1 2X LB : 4) JM101:pINT1 : 2X LB + 0.1 mM IPTG 5) JM101:pINT1 2X LB + 0.5 mM IPTG 6) JM101:pINT1 5X LB 10 7) JM101:pINT1 : 5X LB + 0.1 mM IPTG 8) JM101:pINT1 : 5X LB + 0.5 mM IPTG

Cells were then grown for 24 h at 37°C in a New Brunswick G24 Environmental Incubator Shaker with the shaker speed adjusted to medium setting. At the end of the experiment, the OD₆₀₀ was measured in a Spectronics 21 spectrophotometer by diluting the culture 10-fold in 1% NaC1. The data are listed below.

	<u>Host/Plasmid</u>	LB conc.	IPTG conc. (mM)	Final OD,00
20	HB101:pBR322	2X	. 0	3.00
	JM101:pINT1	2X	0	3.03
	JM101:pINT1	2X .	0.1	2.91
	JM101:pINTl	2X	0.5	3.00
				;
	HB101:pBR322	5X	0	2.73
25	JM101:pINT1	5X	0	3.26
	JM101:pINT1	5X	0.1	3.40
	JM101:pINT1	5X	0.5	3.15

From this data, the following conclusions may be drawn:

30 1. In all cases involving 2X LB, the cells grew to approximately the same density. This density is roughly twice that obtained routinely in 1X LB

under similar growth conditions and indicates exhaustion of available nutrient. In other words, cells have entered stationary phase of growth due to nutrient limitation.

- that cells grown in excess nutrient eventually attain an oxygen-limited growth condition due to which they generate inhibitory metabolics such as acetate.

 Eventually, this leads to cessation of growth, even if more nutrient is supplied. The results of all 5x experiments are indicative of such an occurrence. In other words, oxygen limitation has arisen eventually, causing the culture to reach stationary phase.
- 3. Hence, it may be argued that under
 15 O₂-limited growth, the presence of the hemoglobin
 gene enhances the growth characteristics of <u>E. coli</u>.
 This result is similar to that in Examples 3 and 4,
 with the difference being that here VHb expression is
 not regulated by DO levels.
- 4. It appears that under the given growth conditions, there exists an optimal level of VHb expression that maximizes the growth enhancement effect. Such an optimum may be a function of specific growth properties of each cell line and/or plasmid construct as well as of the environmental conditions of growth. The optimum may thus have to be determined for different applications of this technology on a case-by-case basis; however, such determination does not require undue experimentation.

30 EXAMPLE 7

OF ANOTHER CLONED PROTEIN IN E. COLI.

The aim of this experiment was to demonstrate the effect of the VHb gene on the synthesis of a model cloned gene product. This is an important application of the technology, since a wide variety

of gene products are produced commercially via recombinant DNA technology. A typical process of this kind involves a high cell density fed-batch fermentation. The productivity of such process s is ultimately limited by insufficient oxygen availability.

The following hosts/plasmids were used in this example:

- 1. JM101:pCAT
- 10 2. JM101:pHbCAT.

The construction of these plasmids is described in Example 5.

The following media compositions were used:

LB: 10 g/L Bactotryptone, 5 g/L Yeast Extract,
5 g/L NaC1, 3 g/L K₂HPO₄, 1 g/L KH₂PO₄, 30
mg/L Chloramphenicol

10X feed: 100 g/L Bactotryptone, 100 g/L Yeast Extract, 150 mg/L Chloramphenicol.

The experiment was conducted as follows:

Single colonies of the two strains were inoculated into 5 ml LB in a culture tube and grown overnight. 1 mL of the inoculum was transferred into 100 mL fresh LB and the growth curve was followed. As cells approached the end of log phase, a 1 mL pulse of 10% feed was added and the growth burst was followed. A second pulse was similarly added. At the end of this-growth phase, a pulse of 1 mL 10% feed containing 100 mM IPTG was added to induce the expression of the CAT gene. One hour later, a sample was withdrawn for monitoring CAT activity. The results of the experiment are shown below.

		<u>.</u>	JM101:pCAT	
	1)	Klett before IPTG pulse	e 670	700
	2)	Total soluble protein	0.31	0.435
		(mg/ml culture broth)		
5	3)	CAT activity	1.39x10 ⁴	2.67x104
		(units/mg soluble prote	ein)	
	4)	CAT activity	4.3×10^{3}	11.6×10^{2}
		(units/ml culture broth	a)	

From the above data, the following conclusions may be drawn.

- 1. The presence of VHb enhances the synthesis of a cloned gene product, even at low levels of VHb industrion.
- 2. Besides increasing the amount of cloned
 15 gene product per unit volume of culture, the presence
 of VHb may also enhance the specific activity
 (activity per unit amount of totally soluble protein)
 of the cloned gene product.

EXAMPLE 8

20 OXYGEN-DEPENDENT REGULATION OF
EXPRESSION OF VHb IN E. COLI BY NATIVE
VITREOSCILLA HEMOGLOBIN UPSTREAM SEQUENCES

The aims of this experiment were as follows:

- To demonstrate that VHb gene expression in
 E. coli increases in response to decreasing oxygen levels in the medium.
 - 2. To establish transcriptional-level regulation of gene expression.
- 3. To determine the sensitivity of this oxygen-dependent genetic switch in response to changes in dissolved oxygen concentrations.

Materials and Methods:

The HindIII-SphI fragment containing the VHb gene and flanking sequ nces was cloned into the corresponding sites of the vector pBR322, thereby creating the plasmid pOX1. This was then transformed into the E. coli hold, HB101. The fermentation was conducted in a New Brunswick Bioflo II fermentor with a 2.5 L working volume using LB (10 g/L Bactotryptone, 5 g/L yeast extract, 5 g/L NaC1, 3 g/L K2H-PO4, 1 g/L KH2PO4) plus 8 mg/L silicone antifoam as medium at 37°C, pH 7.0 with a constant agitation speed of 300 rpm. All other methods involve conventional protocols (Maniatis, et al., supra).

Cells were grown to an OD₆₀₀ 0.25 with DO maintained greater than 50% air saturation at all times. At this point, the air supply was gradually reduced so that the DO fell to about 1% air saturation in an almost linear manner over a period of 45 min. (i.e., a time scale long enough for gene induction, yet within approximately one generation time of E. coli). Samples were intermittently taken and analyzed for VHb mRNA and protein levels. Later, nitrogen was sparged in the vessel to study the induction of the VHb promoter under strictly anaerobic conditions.

25 Results:

- 1. The level of VHb mRNA increased about tenfold as DO dropped from 70% to 1% air saturation.
- 2. There was a corresponding increase in VHb activity. A lag was noticed between increase in VHb 30 mRNA level and increase in the quantity of active VHb. This may occur because of the requirement of additional heme biosynthesis in the host cell in order to produce active VHb.
- 3. The VHb promoter was switched on to significant levels only below 40% air saturation and

attains maximum induction levels below 5% air saturation.

4. The promoter switches off under strictly anaerobic conditions, indicating the importance of a basal level of aerobicity in the environment for maximal gene expression.

EXAMPLE 9

OXYGEN-DEPENDENT EXPRESSION OF ANOTHER CLONED PROTEIN

10 In order to use the VHb oxygen-regulated promoter element (ORE) to express other genes, deletions were made from the cloned <u>Vitreoscilla</u> fragment described in Examples 1 and 2 to isolate a functional promoter element. The enzyme Bal 31 was employed for this purpose (Maniatis, <u>et al.</u>, <u>supra</u>). One of the fragments isolated extended from the 5' end of the sequence listed in Example 2 to approximately 125 bp downstream, as sized on a 6% polyacrylamide gel. This fragment was used to express a gene different from that for VHb under oxygen-dependent regulation.

In this experiment, the fragment just described was fused to a promoterless chloramphenicol acetyl transferase (CAT) gene cartridge purchased from Pharmacia Inc. This fusion was inserted into the HindIII-SphI sites of the vector pBR322 to create the plasmid, pOX1. This was transformed into the E. coli host, HB101.

To test the functionality of the ORE, an experiment similar to that in Example 8 was conducted. CAT gene product assays were conducted using a conventional protocol (Neumann, et al., supra).

S. cerevisiae strain 488-0 (leu2, ura3, his 1-7) was transformed with plasmids AAH5 and pEX-2 by the rapid colony transformation procedure (Keszenman-Pereyra and Heida, A colony procedure for transformation of Saccharomyces cerevisiae, Curr. Genet. 13: 21-23, 1988), and plated on synthetic dextrose (SD) medium (Rose, Isolation of genes by complementation in yeast, Methods in Enzymology, 152: 481-504, 1987) without leucine. A representative clonal cell line from each transformation was established after colony purification of a primary transformant.

For the growth studies, single yeast colonies were inoculated into 2 mL of SD -leu (+leu for 488-0) and cultured for 24 hr at 260 rpm at 30°C in a Labline 15 Model 3258 Orbit Enviro-shaker. 0.5 mL of this inoculum was added to 50 mL of the same medium in a 250 mL flask and cultured at 260 rpm at 30°C. Cell growth was measured by turbidity (A_{500mm}) using a Perkin-Elmer Lambda 4A Spectrophotometer. When the 20 glucose level of the culture medium dropped below 2.0 mM, the cultures were pulsed with 1/40 volume of a concentrated medium containing 20 x SD (40% glucose, 13.3% Difco yeast nitrogen base without amino acids, and 1.6 mg/mL of all the amino acids except leucine. 25 For strain 488-0, 1.6 mg/mL leucine was included in the pulse medium) Glucose concentration was estimated using Ames Glucostix test strips.

Results:

A comparison of the growth curves of strains 488-0, 30 488-0:AAH5, and 488-0:pEX-2 grown under the above conditions revealed the following:

1. All three strains grew at an equivalent rate during the logarithmic stage of growth.

Results:

At DO = 70% air saturation, the CAT activity in

E. coli HB101:pOX1 was 6.1x10⁴ units/mg soluble

prot in. After maintaining the DO between 2 to 5% air

5 saturation for 45 min., CAT activity in E. coli

HB101:pOX1 was 6.3x10⁵ units/mg soluble protein.

This demonstrates the isolation of a functional ORE

capable of expressing proteins other than

Vitreoscilla hemoglobin under control of dissolved

10 oxygen content of the culture.

EXAMPLE 10

GROWTH ENHANCEMENT OF SACCHAROMYCES CEREVISIAE CELLS EXPRESSING VHb

In this example, the effect of VHb expression on the growth of the yeast <u>Saccharomyces cerevisiae</u> was studied. The VHb gene was cloned into a yeast expression plasmid, AAH5, that is stably maintained as an extrachromosomal plasmid in yeast cells.

Materials and Methods:

- Plasmid pEX-2 was constructed as follows. The BamHI/SphI fragment described in Example 5 was cloned by blunt-end ligation into the HindiII site of the yeast expression vector AAH5 (Ammerer, Expression of genes in yeast using the ADC-1 promoter, Methods in
- 25 Enzymology 101: 192-201, 1983). AAH5 contains the selectable yeast marker Leu 2, the 2 micron circle origin of replication, and a unique HindIII site flanked by the transcriptional promoter and terminator regions of the yeast alcohol
- 30 dehydrogenase-1 (ADH-1) gene. The ADH-1 promoter will support high levels of transcription of any sequence cloned into the HindIII site. The ADH-1 gene is constitutively expressed in yeast.

2. The VHb-containing strain 488-0:pEX-2 grew to a final optical density of 13.0, while strains 488-0:AAH5 and 488-0 grew to optical densities of only 10.0 and 9.5, respectively. This represents a 26.0% increase in final cell density b tween a strain carrying the VHb gene on a plasmid (488-0:pEX-2) compared to a strain containing the identical plasmid without the VHb gene (488-0:AAH5). In addition, this represents a 32.6% increase in the final cell density of 488-0:pEX-2 over the strain containing no AAH5-derived plasmid (488-0).

EXAMPLE 11

GROWTH ENHANCEMENT DUE TO EXPRESSION OF VHb IN E. COLI FROM A CHROMOSOMALLY INTEGRATED GENE

15 In this example, the tac-VHb gene fusion, discussed above, was integrated into the chromosome of E. coli MG1655 (obtained from Cold Spring Harbor Laboratory, NY).

Materials and Methods:

- 20 A defective Tn10 transposon (Foster, et al., Three Tn10-associated excision events: Relationship to transposition and role of direct and inverted repeats, Cell, 23: 215-227, 1981) was constructed as follows. A kanamycin resistance gene (Pharmacia
- Inc.) was cloned into the SalI site of pINT1 (Example 5). The EcoRI/EagI fragment from the resulting plasmid, which contains the entire tac-VHb fusion and Kan^R gene, was cloned between the inverted repeats (bases 1-66 on the right end and bases 9234-9300 on
- the left end) of a Tn10 derivative which lacks the transposase gene (obtained from Cold Spring Harbor Laboratory, NY). The resulting element, Tn10dKan-tac-VHb, was cloned into a multicopy plasmid containing a tac-Tn10 rightward transposase (obtained

from Cold Spring Harbor Laboratory, NY). Transposition was induced with 0.5 mM IPTG for 4 hr. f llowing which cells were plated on lactose-MacConk y-Kan plates. Lac mutants were selected and 5 the transposon-induced mutation was induced into E. coli MG1655 using P1 phage (Silhavy, et al., supra). One of the resulting Lac colonies, which was further purified and checked for Lac KanR, Amps, VHb+ (IPTG inducible, as confirmed by assay described in 10 Example 3), was designated GRO13. Comparison of growth properties of strains MG1655 and GR013 in 2X LB (described in Example 6) containing 1 mM IPTG, followed by addition of a concentrated feed (25% Bactotryptone, 12.5% yeast extract), showed an 15 increase in final cell densities (final cell densities: $OD_{600} = 16.8$ for MG1655, $OD_{600} = 18.1$ for GR013).

EXAMPLE 12

Oxygen-regulated Expression of Vitreoscilla

hemoglobin from a single-copy, integrated hemoglobin gene.

Two strains of <u>E</u>. <u>coli</u> were developed that contain the <u>Vitreoscilla</u> hemoglobin gene and its promoter, integrated into the chromosome. The hemoglobin gene region was inserted into the chromosome of wild type strain MG1655 (obtained from the Cold Spring Harbor Laboratories) via transposon-mediated integration. The two strains, GRO21 and GRO22, have the hemoglobin gene region inserted into the host <u>lac</u> and <u>xyl</u> operons, respectively.

In order to determine the extent of oxygen-dependent regulation of the integrated hemoglobin expression was monitored during growth in a shake-flask. The cells were grown in 100 ML of a buffered complex

media [10 g/L yeast extract, 5 g/L yeast extract, g/L NaCl, 3 g/L K₂HPO₄, a g/L KH₂PO₄ (pH 7)] at 200 rpm in a New Brunswick G24 shaker-incubator at 37°C. The larg culture volume ensured that the culture became xygen-limited before any other nutrient became limiting or before significant levels of organic acids had accumulated.

Hemoglobin expression was determined by Western analysis using anti-hemoglobin antiserum. In both strains, hemoglobin was not detectable in early exponential-phase cells when the culture oxygen concentration was high. Later, when oxygen became limited (OD₅₉₀=0.9), hemoglobin levels increased significantly. At an OD₅₉₀ of 2.8, hemoglobin expression decreased, indicating that hemoglobin gene promoter activity is lower when the oxygen concentration is reduced to near-zero levels.

Stability of the integrated genes was confirmed by subculturing the cells and demonstrating hemoglobin expression after twenty generations of growth.

In summary, significant amounts of hemoglobin were expressed from a single, integrated copy of the hemoglobin gene. Activity of the integrated gene was regulated by oxygen and maximal expression was achieved under microaerobic conditions. This is similar to its expression properties on a multicopy plasmid.

EXAMPLE 13

The Vitreoscilla Hemoglobin Gene Promoter Region Contains Two Major Transcriptional Initiation Sites

Primer extension analysis was employed to identify transcriptional start site(s) (promoters) within the upstream region of the <u>Vitreoscilla</u> hemoglobin gene. In order to obtain higher levels of hemoglobin RNA, the <u>Vitreoscilla</u> hemoglobin gene region [Khosla and Bailey (1988) <u>Mol. Gen. Genet.</u>, 214: 158] was cloned into the multicopy plasmid pBR322. The resultant plasmid, pOX1, was then introduced into the <u>E. coli</u> host strain HB101 (FhsdS20 recA13 aral4 proA2 lacY1 galK2 rpsL20 xy15 mtl1 supE44).

A 30-mer oligonucleotide corresponding to bases +32 15 to +1 (relative to the translation initiation codon) of the hemoglobin gene was hybridized to total messenger RNA (mRNA) isolated from HB101pOX1 cells. The oligo/RNA hybrids were extended with reverse transcriptase and radiolabeled trinucleotides 20 [Kingston (1978) in <u>Current Protocols in Molecular</u> Biology, Ausubel, et al., eds., John Wiley, 4.8.1] and the products resolved on an acrylamide/urea sequencing gel. Two major extension products were visualized. The most intense band mapped to position 25 -56 which is located 10 bp downstream of a putative Pribnow box. The less intense band mapped to position -109. This result indicated that there are two major transcription initiation sites within the hemoglobin gene promoter. The identical experiment 30 using mRNA from cells growing at different oxygen concentrations indicated that both promoters are

oxygen-regulated and activated under microaerobic

conditions.

EXAMPLE 14

The Cyclic AMP-Catabolite Activator Pr tein

(cAMP-CAP) Complex is Involved in the Regulation of
the Vitreoscilla Hemoglobin Gene Promoter in

5 Escherichia coli

A plasmid, pOX11, was constructed to study the regulatory properties of the <u>Vitreoscilla</u> hemoglobin promoter in <u>E. coli</u>. pOX11 was constructed by fusing the upstream region and first 11 codons of the <u>Vitreoscilla</u> hemoglobin gene [Khosla and Bailey (1988), <u>Mol. Gen. Genet.</u>, 214: 158], to the <u>E. colilacz</u> gene [Berman, <u>et al.</u>, <u>Gene Anal. Tech.</u>, 1: 43]. The <u>lacz</u> gene contained a small deletion of 5' coding sequence, but retained the ability to synthesize active B-galactosidase (B-gal). This fusion was inserted into the multicopy <u>E. coli</u> plasmid pBR322. B-gal activity was used as the indicator of hemoglobin gene promoter activity.

To test whether the cAMP-CAP system was involved in
the overall control of the promoter, pOX11 was
transformed into a control E. coli strain MC4100
[araD139delta (argF-lac)U169 rpsL150 relA1 flbB5301
ptsF25 deoC1, Casadaban (1986), J. Mol. Biol., 104:
541] and two derivatives of MC4100 and two E. coli
strains that are unable to synthesize either cAMP
[GE1051 (delta cya 854 ilv::Tn10] or CAP [GE1050
(delta crp cam)]. Strains GE1051 and GE1050 were
obtained from Dr. G. E. Weinstock (University of
Texas). In order to achieve maximum induction of the
promoter, the cells were grown in 250 mL shake-flasks
containing 100 mL of buffered complex medium
(described in Example 12) at medium setting in a New
Brunswick G24 shaker-incubator.

Specific B-gal activity was measured in the three strains throughout the growth curve. The maximum Bgal 1 vel observed in the control strain MC4100:pOX11 was above 5000 Miller Units, representing a 10-fold 5 increase over background levels. In contrast, the maximum B-gal levels observed in GE1051:p0X11 were less than 500 Miller units, representing less than a 3-fold increase over background. Addition of cAMP to the GE1051 (cya'):pOX11 culture restored the 10-fold 10 indication seen in the control strain. In addition, nucleotide sequence analysis has identified a consensus CAP binding site at position -91 to -101 (with respect to the translation initiation codon) in the <u>Vitreoscilla</u> hemoglobin gene region. 15 results indicate that the cAMP-CAP complex is involved in the overall regulation of the hemoglobin gene promoter.

EXAMPLE 15

Activation of the Vitreoscilla Hemoglobin Gene 20 Promoter in a Fed-Batch Fermentation

Another plasmid, pOX2, was constructed to study the regulatory properties of the <u>Vitreoscilla</u> hemoglobin gene promoter in <u>E</u>. <u>coli</u>. The plasmid pOX2 was constructed by fusing the promoter to the chloramphenical acetyltransferase (CAT) gene (obtained from Pharmacia). This operon fusion was inserted into the BamHI/HindIII site of the <u>E</u>. <u>coli</u> multicopy plasmid pBR322 to give pOX2. CAT activity [Neuman, <u>et al</u>. (1987), <u>Biotechniques</u>, 5:444] was used as the indicator of promoter activity.

The promoter was introduced into <u>E. coli</u> wild-type strain MG1655 (see Example 12). The promoter activity was monitored throughout a fed-batch growth experiment where the culture dissolved oxygen (DO)

was initially kept high and then rapidly dropped. initiate the xperiment, 2.5 mL of a 12-hour seed culture was inoculat d into 2.5 L of a buffered c mplex m dium [2% yeast extract, 0.5% glycerol, 0.3% 5 K₂HPO₄, 0.1% KH₂PO₄, 0.1% trace metal mix (8.3 mM Na_2MoO_4 , 7.6 mM CuSO₄, 8 mM H_3BO_3), 0.1% vitamin mix (0.042% riboflavin, 0.54% pantothenic acid, 0.6% niacin, 0.14% pyridoxine, 0.006% biotin, 0.004% folic acid), 1 mM MgSO4, 0.05 mM CaCl2, 0.2mM FeCl3, 50 10 μ g/mL ampicillin, pH 7) in a New Brunswick Bioflo III fermentor. The air flow rate was kept constant at 4L/minute. The DO was maintained above 50% of air saturation by increasing the agitation rate as the cells grew. When the culture reached an OD590 of 2.5, 15 concentrated feed medium [43% glycerol, 11% (NH₄)₂SO₄, 0.8% MgSO4, 1% trace metal mix, 1% vitamin mix, 0.2 mM FeCl₃, 0.05 mM CaCl₂, and 40 μ g/ml ampicillin] and 50% yeast extract (added separately) were added at a rate of 3 mL/hour. The feed rates were progressively 20 increased to maintain exponential growth until the culture reached on OD500 of 12.0. During this time, the specific CAT activity increased from 2.1 units/mL-OD₅₉₀ to 5.9 units/mL-OD₅₉₀, indicating a 3fold induction of the promoter occurred at high DO 25 levels.

When the cells reached an OD₅₉₀ of 12.0, the air supply was reduced to 3 L/minute and the agitation lowered to 500 rpm. Under these conditions, the culture DO dropped to 0-1% air saturation within one minute. Within 45 minutes, the CAT activity increased to a maximum of 64.3 units/mL-OD₅₉₀. This indicated an 11-fold induction of the promoter occurred by reducing the DO.

These results indicate that a reduction in culture oxygen concentration can substantially activate the <u>Vitreoscilla</u> hem globin gene promoter in a fed-batch fermentation. In addition, there is at least one oth r regulator of promoter activity besides oxygen concentration. In this example, a greater than 30-fold overall induction of CAT was observed.

EXAMPLE 16

The Early, Oxygen-Independent Induction of the

Vitreoscilla Hemoglobin Gene Promoter is Suppressed
by High Levels of Yeast Extract

The experiment in Example 15 indicated that there is at least one additional mode of the promoter regulation other than oxygen concentration. This 15 regulation was apparent in exponential-phase cells when the DO was kept above 50% air saturation in a fed-batch fermentation. To address whether a culture medium component was involved in the oxygenindependent regulation, MG1655:pOX2 cells were grown 20 in a fed-batch fermentation with different concentrations of yeast extract. The batch and feed media were the same as in Example 15, except that either 0.1%, 0.5%, or 2.0% (w/v) yeast extract was used. There was no yeast extract in the feed medium. 25 The cells were grown in batch mode until an OD of approximately 2.5. Feed medium was then introduced at a rate of 2.5 mL/hour. By adjusting the air flow and agitation rates, the DO was maintained above 50% air saturation for the entire experiment. CAT

30 activity was monitored throughout the experiment. The maximum values as follows:

	Yeast Extract (%)	<u>OD590</u>	CAT (units/mL-OD590)
	0.1%	4.0	11432
	0.5%	4.0	4544
35	2.0%	4.0	1606

5

These results indicate that the early oxygenindependent induction of the <u>Vitreoscilla</u> hemoglobin gene promoter is suppressed by high levels of yeast extract.

EXAMPLE 17

Enhancement of Cloned Chloramphenicol
Acetyltransferase Production in a Fed-Batch
Fermentation Using E. Coli Cells Expressing
Hemoglobin from a Single-Copy, Integrated gene

- 10 A plasmid containing the chloramphenicol acetyltransferase (CAT) gene was introduced into E. Coli strains GRO22 and MG1655 (see Example 12) in order to examine the effect of hemoglobin expression on the production of a cloned protein. The plasmid, pTCAT, was constructed by inserting the CAT gene (obtained from Pharmacia) into the multicopy plasmid pKK223-3 (obtained from Pharmacia) which places the CAT gene under control of the tac (trp-lac) promoter. The tac promoter is constitutively active in MG1655.
- The cells were grown in a 2.5 L fermentor in a buffered defined medium containing 0.5% glucose, 0.15% KH₂PO₄, 0.43% K₂HPO₄, 0.04% (NH₄)₂SO₄, 0.05 mM CaCl₂, 0.2 mM FeCl₃, 1 mM MgSO₄, 0.1% trace metals (Example 15), 0.1% vitamins (Example 15), and 50 μg/mL ampicillin. The agitation speed was maintained at 300 rpm and the air flow rate maintained at 0.2 L/minute. When the culture reached an OD₅₉₀ of 1.2, feed medium (Example 15, except that 43% glucose was used instead of glycerol) was added at a rate of 1.4 mL/hour. The feed rate was increased to 2.8 mL/hour when the culture reached an OD₅₉₀ of 2.2.

CAT activity was monitored throughout the growth period. The results are shown in the following Table:

	<u>Strain</u>	Time (h)	CAT Activity (units/mL)	L
5	MG1655:pTCAT	10	330	
	MG1655:pTCAT	12	380	
	MG1655:pTCAT	14	440	
	MG1655:pTCAT	15	480	
	MG1655:pTCAT	21	430	
		•		
10	GRO22: pTCAT	11	420	
	GRO22:pTCAT	14	740	
	GRO22:pTCAT	15	820	
	GRO22:pTCAT	16	770	

As can be seen, there is a significant enhancement of

15 CAT production in GRO22:pTCAT culminating with a 71%

increase at 15 hours. In addition, there was an

approximately 10% increase in total cell protein in

GRO22:pTCAT cells relative to total cell protein in

MG1655:pTCAT. Western analysis using an anti-CAT

20 antibody indicated that CAT expression was equivalent

in the two strains until they reached oxygen

limitation (approximately 6 hours). This experiment

indicates that an E. coli strain expressing

hemoglobin from a single copy, integrated gene can be

25 used to produce significantly higher quantities of a

cloned protein. In addition, product enhancement

with this system occurs when the culture oxygen

concentration is low.

IN THE CLAIMS:

- 1. A method for preparing any foreign protein in a host cultured cell which grows within a first level f environmental oxygen comprising:
- DNA expression vector capable of replicating in said host cultured cell, said vector having a promoter/regulator capable of directing transcription and translation initiation and control of foreign recombinant-DNA sequences for coding for said foreign protein located downstream therefrom, said promoter/regulator being derived from an organism other than said host, and said promoter/regulator further being capable of being activated upon a decrease of environmental oxygen to below said first level;
 - (b) culturing said host cultured cell in a medium appropriate for expressing the protein expressed by said foreign DNA; and
- (c) lowering the level of oxygen available to said medium to a second level below said first level, said second level providing an oxygen concentration which activates said promoter/regulator.
- A method for preparing any foreign protein in a
 host cultured cell which grows within a first level of environmental oxygen comprising:
- (a) introducing into said host cultured cell a
 DNA expression vector comprising a series of
 nucleotides including the <u>Vitreoscilla</u> hemoglobin

 30 transcription and/or translation initiation sequences
 capable of directing intracellular production of a
 major portion of the <u>Vitreoscilla</u> hemoglobin protein,
 said vector containing foreign DNA sequences coding
 for said foreign protein, said vector having a

 35 promoter/regulator capable of being activated by a

d crease of environmental oxygen to below said first level:

- (b) growing said host cultured cell in a medium appropriate for isolating the protein expressed by5 said foreign DNA; and
 - (c) lowering the level of oxygen available to said medium to a second level below said first level, said second level providing an oxygen concentration which activates said promoter/ regulator.
- 10 3. The method of Claim 2 wherein said portable DNA sequence is introduced directly and integrated into the chromosome of a host cultured cell.
- 4. A method for increasing the growth characteristics, including the growth yield, the growth rate, and the achievable cell density under controlled circumstances, of a host cultured cell grown in the presence of oxygen, said host cultured cell being chosen and derived from a selection of cultured cells consisting of microorganisms and cells obtained from multi-cellular organisms selected from the group consisting of animals, plants and insects comprising:
- (a) preparing a portable DNA sequence capableof directing said host cultured cell to produce aprotein having at least some oxygen-binding activity;
 - (b) introducing said portable DNA sequence into a host cultured cell capable of expressing at least some oxygen-binding protein;
- (c) culturing the host cell under conditions30 appropriate for expression of the protein; and
 - (d) permitting the protein to assume an active structure whereby it possesses at least some oxygen-binding activity.

- 5. The method of Claim 4 wherein said portable DNA sequence is introduced directly and int grated into the chromosome of a host cultured cell.
- 6. The method of Claim 4, wherein said portable DNA5 sequence is introduced into said host cultured cell by the following steps:
- (a) cloning the portable DNA sequence into a vector capable of being transferred into and replicating in a host cell, such vector containing
 operational elements for the portable DNA sequence;
 - (b) transferring the vector containing the portable DNA sequence and operational elements into a host cultured cell capable of expressing at least some of the oxygen-binding protein; and
- 15 (c) culturing the host cell under conditions appropriate for replication and propagation of the vector and expression of the protein.
- A method for increasing the production of proteins, both those normally made and those
 expressed as a result of genetic engineering, biopolymers, and other metabolic products, of a host cultured cell grown in the presence of oxygen, said host cultured cell being chosen and derived from a selection of cultured cells consisting of
- 25 microorganisms and cells obtained from multi-cellular organisms selected from the group consisting of animals, plants and insects comprising:
- (a) preparing a portable DNA sequence capable of directing a host cultured cell to produce a
 30 protein having oxygen-binding activity;
 - (b) introducing said portable DNA sequence into a host cultured cell capable of expressing at least some oxygen-binding protein;
- (c) culturing the host cell under conditions appropriate for expression of the protein; and

- (d) permitting the protein to assume an active structure whereby it possesses at least some oxyg n-binding activity.
- 8. A method for transporting and supplying oxygen5 to oxygen-requiring processes and operations,comprising:
 - (a) preparing a portable DNA sequence capable of directing a host cultured cell to produce a protein having oxygen-binding activity;
- 10 (b) introducing said portable DNA sequence into a host cultured cell capable of expressing at least some oxygen-binding protein;
 - (c) culturing the host cell under conditions appropriate for expression of the protein;
- 15 (d) permitting the protein to assume an active structure whereby it possesses at least some oxygen-binding activity; and
- (e) effectively delivering said host cell or a preparation from said host cell containing the20 oxygen-binding protein to said oxygen-requiring process.
 - 9. A method for the binding and removal of oxygen form an environment comprising:
- (a) preparing a portable DNA sequence capable 25 of directing a host cultured cell to produce a protein having oxygen-binding activity;
 - (b) introducing said portable DNA sequence into a host cultured cell capable of expressing at least some oxygen-binding protein;
- 30 (c) culturing the host cell under conditions appropriate for expression of the protein;
 - (d) permitting the protein to assume an active, structure whereby it possesses at least some oxygenbinding activity; and

- () effectively delivering said host cell or a preparation from said host cell containing the oxygen-binding protein to said oxygen-containing environment.
- 5 10. A method for increasing the growth characteristics, including the growth yield, the growth rate, and the achievable cell density under controlled circumstances, of a cell in culture, said host cultured cell being chosen and derivatized from a selection of cultured cells consisting of microorganisms and cells obtained from multi-cellular organisms selected from the group consisting of animals, plants, and insects, said cell being capable of producing a protein having at least some oxygen-binding activity, comprising:
 - (a) culturing said host cell under conditions appropriate for expression of said protein, and
- (b) permitting said protein to assume an active structure whereby it possesses at least some oxygen-20 binding activity.
- 11. A method for increasing the production of proteins, both those normally made and those expressed as a result of genetic engineering, biopolymers, and other metabolic products, of a host cultured cell grown in the presence of oxygen, said host cultured cell being chosen and derived from a selection of cultured cells consisting of microorganisms and cells obtained from multi-cellular organisms selected from the group consisting of animals, plants and insects, said cell being capable of producing a protein having at least some oxygen-binding activity, comprising:
 - (a) culturing said host cell under conditions appropriate for expression of said protein, and

- (b) permitting said pr tein to assume an active structure whereby it possesses at least some oxygen-binding activity.
- 12. A method according to any one of the Claims 1
 5 through 30 further comprising the step of modulating the activity of said promoter/regulator by control of the amount of cAMP-CAP complex in said host cell.
- 13. A method according to Claim 12 wherein said step of modulating said activity comprises introducing
 10 cAMP into said host cell, wherein said host cell is unable to synthesize CAP or cAMP.
 - 14. A method according to Claim 12 wherein said step of modulating said activity comprises manipulation of the activity of a crp gene in said host.
- 15 15. A method according to any one of Claims 4 through 11 further comprising the step of modulating the expression of said protein by control of the amount of cAMP-CAP complex in said host cell.
- 16. A method according to Claim 15 wherein said step
 20 of modulating said expression comprises introducing
 cAMP into said host cell, wherein said host cell is
 unable to synthesize CAP or cAMP.
- 17. A method according to Claim 16 wherein said step of modulating said expression comprises manipulation25 of the activity of a crp gene in said host.
 - 18. A method for regulating the production of at least one designated protein, selected from the group consisting of those proteins normally made and those expressed as a result of genetic engineering,
- 30 biopolymers, and other metabolic products, of a host

cultured cell grown in the presence of oxygen, said
host cultured cell being chosen and derived from a
selection of cultured cells c nsisting of
microorganisms and cells obtained from multi-cellular
organisms selected from the group consisting of
animals, plants and insects comprising:

- (a) introducing a portable DNA sequence capable of directing a host cultured cell to produce a protein having oxygen-binding activity into a host 10 cultured cell;
 - (b) culturing said host cell under conditions appropriate for expression of said oxygen-binding protein and said designated protein.
- 19. A method according to Claim 18 wherein said15 designated protein is expressed from an extrachromosomal element in said cell.
 - 20. A method according to Claim 19 wherein said oxygen-binding protein comprises <u>Vitreoscilla</u> hemoglobin.
- 20 21. A method according to Claim 20 wherein said designated protein is chloramphenicol acetyltransferase.
- 22. A method for preparing any foreign protein in a host cultured cell which grows within a first level 25 of environmental oxygen comprising:
- (a) introducing into said host cultured cell a
 DNA expression vector capable of replicating in said
 host cultured cell, said vector having a
 promoter/regulator capable of directing transcription
 and translation initiation and control of foreign
 recombinant-DNA sequences coding for said foreign
 protein located downstream therefrom, said
 promoter/regulator being derived from an organism

other than said host, and said promoter/regulator being capable of being activated upon a decrease of environmental oxygen to below said first level and further capable of being suppressed ind pendently of environmental oxygen concentration during exponential growth of said host cultured cell;

- (b) culturing said, host cultured cell in a suitable medium appropriate for expression of said foreign DNA; and
- (c) introducing a source of complex nitrogen into said medium during exponential growth of said host to a level sufficient to suppress said promoter/regulator.
- 23. A method of preparing any foreign protein in a
 15 host cultured cell which grows within a first level
 of environmental oxygen comprising:
- (a) introducing into said host cultured cell a
 DNA expression vector comprising a portable DNA
 sequence comprising a series of nucleotides capable
 of directing intracellular production of a major
 portion of the <u>Vitreoscilla</u> hemoglobin protein, said
 vector containing foreign DNA sequences coding for
 said foreign protein, said vector having a
 promoter/regulator capable of being activated by a
 decrease of environmental oxygen to below said first
 level and capable of being suppressed independently
 of environmental oxygen concentration during
 exponential growth of said host cultured cell;
- (b) growing said host cultured cell in a medium30 appropriate for expression of said foreignDNA; and
- (c) introducing a source of complex nitrogen into said medium during exponential growth of said host to a level sufficient to suppress said 35 promoter/regulator.

- 24. The method of Claim 23, wherein said portable DNA sequence is introduced directly and integrated into the chromosome of a host cultured cell.
- 25. The method of Claim 23 wherein said portable DNA 5 sequence comprises the <u>Vitreoscilla</u> hemoglobin transcription and/or translation initiation sequences.
 - 26. The method according to Claim 22 wherein said source of complex nitrogen comprises yeast extract.
- 10 27. The method according to Claim 23 wherein said source of complex nitrogen comprises yeast extract.

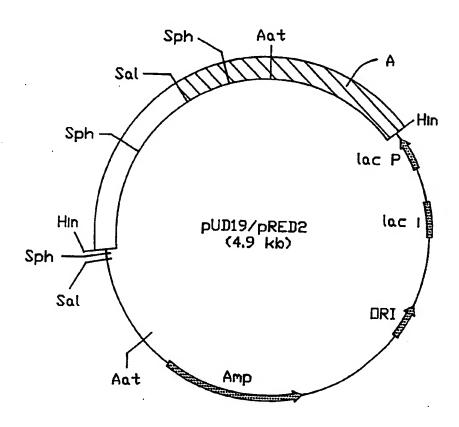


FIG.-1

SUBSTITUTE SHEET

INTERNATIONAL SEARCH REPORT

	IN TERRET TOTAL	SEARCH REPURI			
International Application No PCT/US90/06083 I. CLASSIFICATION F SUBJECT MATTER (if several classification, symbols apply, indicate all) 3					
According to International Patent Classification (IPC) or to both National Classification and IPC					
IPC(5)): C12N 15/00, 15/31, 1/00; A611	29/01			
<u>U.S.C1</u>	: 435/69.1, 172.3, 317 _. 1, 266				
II. FIELD	S SEARCHED				
	Minimum Docume	ntation Searched •			
Classificati	on System	Classification Symbols			
บ.ร. c	U.S. C1. 435/69.1, 172.3, 317.1, 266, 42, 801, 818				
	Documentation Searched other to the Extent that such Occuments	than Minimum Documentation are Included in the Fields Searched •			
ļ	See Attachment				
III. DOCL	JMENTS CONSIDERED TO BE RELEVANT :+				
Category *	<u> </u>				
XY	WO, A, 89/03883 (Khos)	la et al.) 05 May 1989,			
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	in <u>Escherichia coli</u> ", entire document.	pages 158-161. See			
	entire document.	;			
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	d categories of cited documents: 13	"T" later document published after the international filing date			
	"A" document defining the general state of the art which is not considered to be of particular relevance or pnority date and not in conflict with the application but cited to understand the principle or theory underlying the				
"E" earl	ler document but published on or after the International o	invention "X" document of particular relevance; the claimed invention			
"L" document which may throw doubts on priority claim(s) or cannot be considered novel or cannot be considered to involve an inventive step.					
cita	which is cited to establish the publication date of another citation or other special reason (as specified) "Y" document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the				
"O" document referring to an oral disclosure, use, exhibition or other means document is combined with one or more other such documents, such combination being obvious to a person skilled					
"P" document published prior to the international filing date but later than the priority date claimed "a" document member of the same patent family					
IV. CERTIFICATION					
Date of the Actual Completion of the International Search 2 Date of Mailing of this International Search Report 2					
16 January 1991					
14 January 1991					
Internation	al Searching Authority 1	Signature of Authorized Officer 10			

Form PCT/ISA/210 (second sheet) (May 1988)

III. DOCUM	ENTS CONSIDERED TO BE RELEVANT (CONTINUED FROM THE SECOND S	CT/US90/06083
Category •	Citation of Document, 10 with indication, where appropriate, of the relevant passages 17	
X	Nature, volume 331, issued 18 February Khosla et al, "Heterologous expression bacterial haemoglobin improves the gropperties of recombinant Escherichia opages 633-635. See entire document.	offa 18-20 wth
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International Application No. PCT/US90/06083		
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	S WHERE CERTAIN CLAIMS WERE FOUND UNSEARCHABI	·
	report has not been established in respect of certain claims under Ar	
1. Claim numbers	, because they relate to subject matter I not required to be search	ed by this Authority, namely:
2. Claim numbers	, because they relate to parts of the international application that	do not comply with the prescribed require
ments to such an ex	extent that no meaningful international search can be carried out 1, specified	ecifically:
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3. Ctaim numbers	, because they are dependent claims not drafted in accordance w	uth the second and third sentances of
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VI. OBSERVATION	S WHERE UNITY OF INVENTION IS LACKING ²	
This International Search	ting Authority found multiple inventions in this international application	on as follows:
	hment to Telephone practice	
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1.X As all required addit	itional search fees were timely paid by the applicant, this international	search report covers all searchable claim
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Form PCT/ISA/210 (supplemental sheet (2) (Rev. 4-90)

Attachment To PCT/IPEA/210 USPTO Automated Patent System (file USPAT, 1970-1991), CAS.

Search terms:

Vitreoscilla, hemoglobin, globin, artificial blood, oxygen transport; author search Khosla, C; Bailey, J; Dikshit, K; Webster, D.